SA 101429

## INTERNATIONAL SEARCH REPORT

International application No.

25/02/95

PCT/US 94/14073

	locument arch report	Publication date		family nber(s)	Publication date
WO-A1-	9308279	29/04/93	AU-A-	2794092	21/05/93
			AU-A-	2865492	21/05/93
			CA-A-	2121497	29/04/93
			CA-A-	2121498	29/04/93
			EP-A-	0610335	17/08/94
			EP-A-	0623168	09/11/94
			FI-A,D-	941757	16/06/94
			FI-A,D-	941758	16/06/94
			HU-D-	9401108	00/00/00
•			HU-D-	9401109	00/00/00
			NO-A.D-	941370	08/06/94
			NO-A.D-	941371	15/06/94
-	*	•	PT-A-	100978	31/01/94
			PT-A-	100979	31/01/94
			WO-A-	9308280	29/04/93

#### INTERNATIONAL SEARCH REPORT

CONTRACTOR OF THE PROPERTY OF

International application No. PCT/US 94/14073

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Box I	Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)
This inte	mational search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
1. X	Claims Nos.: 37 because they relate to subject matter not required to be searched by this Authority, namely:
	See PCT Rules 39 (iv): Method for treatment of the human body by therapy.
	·
2.	Clairns Nos.:
	because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3.	Claims Nos.: because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II	Observations where unity of invention is lacking (Continuation of Item 2 of first sheet)
This Inte	rnational Searching Authority found multiple inventions in this international application, as follows:
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1.	As all required additional search sees were timely paid by the applicant, this international search report covers all searchable claims.
2.	As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3.	As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4.	No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:
	: ·
Remark	The additional search fees were accompanied by the applicant's protest.  No protest accompanied the payment of additional search fees.

## INTERNATIONAL SEARCH REPORT

International application No. PCT/US 94/14073

C (Continu		Relevant to claim No.
A	WO, A1, 9308279 (IMMULOGIC PHARMACEUTICAL CORPORATION), 29 April 1993 (29.04.93)	1-36,38-47
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#### INTERNATIONAL SEARCH REPORT

Inte. .tional application No. PCT/US 94/14073

#### A. CLASSIFICATION OF SUBJECT MATTER IPC6: C07K 14/435 According to International Patent Classification (IPC) or to both national classification and IPC B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) IPC6: C07K Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practicable, search terms used) MEDLINE, BIOSIS, DERWENT BIOTECHNOLOGICAL ABSTRACTS, WPI, CLAIMS C. DOCUMENTS CONSIDERED TO BE RELEVANT Category\* Citation of document, with indication, where appropriate, of the relevant passages Relevant to claim N . X IMMUNOLOGY, Volume 75, 1992, G. A. STEWART et al, 1-36,38-47 "The group III allergen from the house dust mite Dermatophagoides pteronyssinus is a trypsin-like enzyme", page 29 - page 35, summary, and page 34, column 1, lines 44-54 P,X CLINICAL AND EXPERIMENTAL ALLERGY, Volume 24, 1994, W.-A. SMITH et al, "Cloning and sequencing of the 1-36,38-47 Dermatophagoides pteronyssinus group III allergen, Der p III", page 220 - page 228, the whole document X Further documents are listed in the continuation of Box C. See patent family annex. Special categories of cited documents later document published after the international filing date or priority document defining the general state of the art which is not considered. date and not in conflict with the application but cited to understand the principle or theory underlying the invention to be of particular relevance ·Eertier document but published on or after the international filing date document of particular relevance: the claimed invention cannot be considered novel or cannot be considered to involve an inventive document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other step when the document is taken alone special reason (as specified) document of particular relevance: the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination "O" document referring to an oral disclosure, use, exhibition or other means. being obvious to a person skilled in the art document published prior to the international filing date but later than the priority date claimed "&" document member of the same patent family Date of the actual completion of the international search Date of mailing of the international search report 2 5. 04. 95 29 March 1995 arne and mailing address of the International Searching Authorit Authorized officer European Parent Office, P.B. 5813 Patentiaan 2 NL 2280 HV Rijsmik PATRICK ANDERSSON Tel. (+31-70) 340-2040, Tx. 31 651 epo ni. Fax: (+31-70) 340-3016

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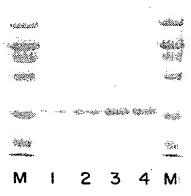


Fig. 3

Derpii Crayfish	AEVTDNMICG DEIFDSMICA	GDVANGGKDS G-VPEGGKDS	COGDSGGPVV COGDSGGPLA	DVKNNQVV ASDIGSTYLA	AEVIDNMICG GDVANGGKDS CQGDSGGPVVDVKNNQVV GIVSWGYGCA DEIFDSMICA G-VPEGGKDS CQGDSGGPLA ASDIGSTYLA GIVSWGYGCA	237
Consensus	E.D.MIC.	D.MIC. G.V GGKDS CQGDSGGPD	COGDSGGP	g.	GIVSWGYGCA	250
Derpii CRAYFISH	RKGYPGVYTR RPGYPGVYTE	RKGYPGVYTR VGNFIDWIES KRSQ RPGYPGVYTE VSYHVDWIKA N-AV	KRSQ N-AV			261 237
Consensus	R.GYPGVYT.	R.GYPGVYI. VDWI	•			274

Derpii CRAYFISH	MITYNILIVL	IYNILIVL LLAINTLANP	Ilpaspnati I	ILPASPNATI VGGEKALAGE	CPYQISLQS- FPYQLSFQET	4 2
Consensus	•	•	-	I VGGAGE .PYQ.S.Q	.PYQ.S.Q	20
Derpii	FLGFSFHFCG	GTILDEYWIL ASIYNENYAI	Taahcvag Taghcvygdd	-QTASKLSIR YENPSGLQIV	Ynslkhsl Ageldmsvne	9.5
Consensus	S.HFCG	S.HFCGIE	TA.HCV.G	S.L.H.	r.s	Ä
Derpii CRAYFISH	GGEK-ISVAK IFAHEKYDSY GSEQTITVSK IILHENFDYD	IFAHEKYDSY IILHENFDYD	QIDNDIALIK LLDNDISLLK	QIDNDIALIK LKSPMKLNOK NAKAVGLPAK LLDNDISLLK LSGSLTFN-N NVAPIALPAQ	nakavglpak Nvap ialpaq	ਜੋ ਜੋ
Consensus	G.EI.V.K	E.I.V.K IHED	DNDI.L.K	DNDI.L.K LN. NLPA.	NLPA.	ਜ
Derpii CRAYFISH	GSDVKVGDQV G-HTATGN-V	RVSGWGYLEE IVTGWGTTSE		GSYSLPSELR RVDIAVVSRK ECNELYSKAN GG-NTPDVLQ KVTVPLVSDA ECRDDYGA	ECNELYSKAN ECRDDYGA	ਜੌਜ
Consensus	GG.`V	GV .V.GWGE GPLVVS ECY	GPL.	.vvs	EC Y	Ñ

602	662	722	782	842
AGTTATTCATTACCATCTGAATTAAGACGTGTTGATATTGCTGTTATCACGTAAAGAA 602 S Y S L P S E L R R V D I A V V S R K E 140	TGTAATGAATTATATTCAAAAGCTAATGCTGAAGTTACTGATATGATTTGTGGTGGT 662 C N E L Y S K A N A E V T D N M I C G G 170	GATGTTGCAAATGGTGGTAAAGATTCTTGTTGATTCTGGTGGACCGGTTGTTGAT 722 DVANGGKDSCTGTTGTCAAGGTGATTCTGGTGGACCGGTTGTTGAT 722 DVANGG KDSCQG GDSGG PVVD 180	GITAAAATAATCAAGTTGTTGTTTCATGGGGTTATGGTTGTGCACGTAAAGGT 782 V K N N Q V V G I V S W G Y G C A R K G 200	TATCCAGGTGTTTATACACGTGTAGTTTGAATCAAACGTTCA 842 Y P G V Y T R V G N F I D W I E S K R S 220
AGTTATT S Y	TGTAATG C N	GATGTTG D V	GITAAAA V K 1	TATCCAG( Y P (

1022 902 CAGTGATTGATAAAACAATTTCGACATATAAAAATTGGCAAATGATGCCATTATATGTT

122	182	242	302	362	422	482	542
TITITITITITICAAAACAITGAAAACATTGAAGCAACTGTGTGTACAACCAGAAAG ATGATCATCTATAATATTTTATTT	CTACC L E	TGTCCATATCAGTTACAATCAAGTAGTCATTTTTGTGGTGGTACTATTCTTGAT CPYQISSSHFCGGTGTTTTTGTGTGTACTATTCTTGAT 20	GAATATTGGATTTTAACAGCTGCACTTGTGTTGCCGGACAAACAGTAAACTTTCA E Y W I L T A A H C V A G Q T A S K L S E Y W I L T A A H C V A G Q T A S K L S	ATTCGTTACAATAGTTTAAAATTTAGGTGGTGAAAAATTTCTGTTGCTAAAATT I R Y N S L K H S L G G E K I S V A K I 60	TITGCACATGAAAATATGATAGTTATCAAATTGATAATGATAAGCTT F A H E K Y D S Y Q I D N D I A L I K L 80	AAATCACCTATGAAATTCAGAAAATGCCAAAAGCTGTTGGATTACCAGCAAAAGGA KSPMKLNQKNAA KAVGLPPAKG 100	TCGGATGTAAAAGTTGGTGATCGAGTTCTGGTTAGGGAAGGAA

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- 35. A composition of claim 34, wherein the peptide comprises an amino acid sequence of Figure 1A and 1B (SEQ ID NO:2).
- 36. A composition of claim 35, wherein the peptide comprises amino acid residues 1-232 of the amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2).
  - 37. A method of treating sensitivity to a house dust mite allergen in a subject sensitive to the allergen, comprising administering to the subject the composition of claim 34.
  - 38. A method of detecting sensitivity in a subject to a house dust mite allergen, comprising combining a blood sample obtained from the subject with a peptide of claim 22, under conditions appropriate for binding of blood components with the peptide and determining the extent to which such binding occurs.
  - 39. A method of claim 38, wherein the extent to which binding occurs is determined by assessing T cell function, T cell proliferation, B cell function, binding of the protein to antibodies present in the blood or a combination thereof.
    - 40. An antibody specifically reactive with a peptide of claim 22.
    - 41. An antibody of claim 40 which is a monoclonal antibody.
- 25 42. An antibody specifically reactive with a peptide of claim 25.
  - 43. An antibody of claim 42 which is a monoclonal antibody.
  - 44. A T cell clone specifically reactive with a peptide of claim 22.
  - 45. A soluble T cell receptor specifically reactive with a peptide of claim 22.
  - 46. An antibody specifically reactive with the T cell receptor of claim 45.
- The antibody of claim 46 wherein the antibody is a monoclonal antibody.

- 24. An isolated peptide having an activity of a house dust mite allergen, *Der p* III, produced by recombinant expression of a nucleic acid of claim 4.
- 25. An isolated peptide having an activity of a house dust mite allergen, *Der p* III, produced by recombinant expression of a DNA of claim 11.
- 26. An isolated peptide having an activity of a house dust mite allergen, *Der p* III, produced by chemical synthesis.
- 27. An isolated peptide of claim 26 which is at least about 10-20 amino acid residues in length.
- 28. An isolated peptide of claim 26 which is at least about 10-16 amino acid residues in length.
  - 29. A modified peptide having an activity of Der p III.
- 30. A modified peptide of claim 29, wherein at least one cysteine residue present in the *Der p* III amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2) is replaced by another amino acid.
- 31. A modified peptide of claim 30, wherein at least one cysteine residue present in the *Der p* III amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2) is replaced by a serine residue.
  - 32. A modified peptide of claim 29, wherein at least one lysine residue is added to either the amino or carboxy terminus or both the amino and carboxy terminus of the peptide.
  - 33. A substantially pure preparation of a peptide having an activity of a house dust mite allergen, *Der p* III.
    - 34. A composition suitable for pharmaceutical administration comprising at least one peptide having an activity of *Der p* III and a pharmaceutically acceptable carrier.

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- 11. An isolated DNA comprising a nucleotide sequence encoding a peptide of at least about 10-16 amino acid residues or more in length and having at least about 50% homology with an amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2).
  - 12. A recombinant expression vector comprising the nucleic acid of claim 1.
- 13. A recombinant expression vector of claim 12, wherein the nucleic acid is cDNA.
- 10 14. A recombinant expression vector of claim 13, wherein the cDNA comprises the nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1).
  - 15. A host cell transfected with the recombinant expression vector of claim 12 capable of directing the expression of a peptide having an activity of *Der p* III.
    - 16. A host cell of claim 15 which is a eucaryotic cell.
  - 17. A host cell transfected with the recombinant expression vector of claim 13 capable of directing the expression of a peptide having an activity of *Der p* III.
    - 18. A host cell of claim 17 which is a eucaryotic cell.
  - 19. A host cell transfected with the recombinant expression vector of claim 14 capable of directing the expression of a peptide having an activity of *Der p* III.
    - 20. A host cell of claim 19 which is a eucaryotic cell.
- 21. A method of producing a peptide having an activity of *Der p* III comprising culturing a host cell of claim 15 in medium to express the peptide and isolating the peptide from the culture.
  - 22. An isolated peptide having an activity of a house dust mite allergen, *Der p* III. produced by recombinant expression of a nucleic acid of claim 1.
- 35 23. An isolated peptide having an activity of a house dust mite allergen, *Der p* III, produced by recombinant expression of a nucleic acid of claim 3.

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### What is claimed is:

- 1. An isolated nucleic acid comprising a nucleotide sequence encoding a peptide having an activity of a house dust mite allergen, *Der p* III.
  - 2. An isolated nucleic acid of claim 1, which is a cDNA sequence.
- 3. An isolated nucleic acid of claim 2, wherein the cDNA comprises a nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1).
- 4. An isolated nucleic acid of claim 2, wherein the cDNA comprises the coding region of a nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1).
- 5. An isolated nucleic acid of claim 1, wherein the peptide comprises an amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2).
  - 6. An isolated nucleic acid of claim 5, wherein the peptide comprises amino acid residues 1-232 of the sequence shown in Figure 1A and 1B (SEQ ID NO:2).
- 7. An isolated nucleic acid of claim 1, wherein the peptide is at least 50% homologous with a sequence comprising an amino acid sequence shown in Figure 1A and 1B (SEO ID NO:2).
- 8. An isolated nucleic acid of claim 1, wherein the peptide is encoded by a nucleic acid which hybridizes under high or low stringency conditions to a nucleic acid which encodes a peptide comprising an amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2).
- 9. An isolated nucleic acid of claim 1, wherein the peptide is at least about 10-20 amino acid residues in length.
  - 10. An isolated nucleic acid of claim 1, wherein the peptide is at least about 10-16 amino acid residues in length.

	(xi)	SEQU	JENCI	E DES	SCRIE	OIT	1: S	EQ II	NO:	:16:						
	Ile 1	Val	Gly	Gly	Glu 5	Lys	Ala	Leu	Ala	Gly 10	Gln	Ser	Pro	Tyr	Gln 15	Ile
	Ser	Leu	Gln	Ser 20	Ser	Ser	His	Phe	Ser 25	Gly	Gly	Thr	Ile	Leu 30		
(2)	INFO	RMAT:	ION 1	FOR S	SEQ I	ID NO	0:17	:								
	(i)	(A)	LEI	NGTH PE: 8	: 10 amino	amin ac:	no a id			,						
	(ii)		•													
			•						÷							
	(xi)	SEQ	UENC	E DE	SCRI	PTIO	N: S	EQ I	D NO	:17:						
	Met 1	Gly	His	His	His 5	His	His	His	Glu	Phe 10						
(2)	INFO	RMAT	ION	FOR	SEQ	ID N	0:18	:								
	(i)	(A (B (C	) LE ) TY ) ST	NGTH PE: RAND	: 20 nucl EDNE	bas eic SS:	e pa acid sing	irs			-	s.	-			
	(ii)	MOL	ECUL	E TY	PE:	CDNA										
	(xi)	SEQ	UENC	E DE	SCRI	PTIO	N: S	EQ I	D NO	:18:						
GCG	GATCC	י דים: "מי	GTTG	GTGG	;T											
	(2)	Ile 1 Ser (2) INFO (ii) (xi) Met 1 (2) INFO (ii) (iii) (xii)	Ile Val 1 Ser Leu  (i) SEQU (A) (B) (D) (ii) MOLI (v) FRAC (xi) SEQU Met Gly 1  (2) INFORMAT (i) SEQU (A) (B) (C) (A) (A) (B) (C) (A) (B) (C) (A) (B) (C) (A) (B) (C) (B) (C) (C) (C) (C) (C) (C) (C) (C) (C) (C	Ile Val Gly 1 Ser Leu Gln  (i) SEQUENCY (A) LEY (B) TYY (D) TOY  (ii) MOLECULY (v) FRAGMENT  (xi) SEQUENCY Met Gly His 1  (2) INFORMATION  (i) SEQUENCY (A) LEY (B) TYY (C) ST (D) TOY  (ii) MOLECULY (xi) SEQUENCY (xi) SEQUENCY	Ile Val Gly Gly  1  Ser Leu Gln Ser 20  (i) SEQUENCE CHE (A) LENGTH (B) TYPE: (D) TOPOLOG  (ii) MOLECULE TY  (v) FRAGMENT TY  (xi) SEQUENCE DE  Met Gly His His 1  (2) INFORMATION FOR (i) SEQUENCE CHE (A) LENGTH (B) TYPE: (C) STRAND (D) TOPOLOG  (ii) MOLECULE TY  (xi) SEQUENCE DE	Ile Val Gly Gly Glu  1 5  Ser Leu Gln Ser Ser 20  (2) INFORMATION FOR SEQ 3  (i) SEQUENCE CHARACT (A) LENGTH: 10 (B) TYPE: amino (D) TOPOLOGY: 3  (ii) MOLECULE TYPE: 1  (v) FRAGMENT TYPE: 3  (xi) SEQUENCE DESCRIPT Met Gly His His His 1  5  (2) INFORMATION FOR SEQ (A) LENGTH: 20 (B) TYPE: nucl (C) STRANDEDNE (D) TOPOLOGY:  (ii) MOLECULE TYPE:	Ile Val Gly Gly Glu Lys  Ser Leu Gln Ser Ser Ser  20  (2) INFORMATION FOR SEQ ID NO  (i) SEQUENCE CHARACTERIS  (A) LENGTH: 10 amin  (B) TYPE: amino acc  (D) TOPOLOGY: lines  (ii) MOLECULE TYPE: pept:  (v) FRAGMENT TYPE: inter  (xi) SEQUENCE DESCRIPTION  Met Gly His His His His I  (i) SEQUENCE CHARACTERI  (A) LENGTH: 20 bas  (B) TYPE: nucleic  (C) STRANDEDNESS:  (D) TOPOLOGY: line  (ii) MOLECULE TYPE: cDNA  (xi) SEQUENCE DESCRIPTION  (xi) SEQUENCE DESCRIPTION	Ile Val Gly Gly Glu Lys Ala  1 5  Ser Leu Gln Ser Ser Ser His 20  (2) INFORMATION FOR SEQ ID NO:17  (i) SEQUENCE CHARACTERISTIC (A) LENGTH: 10 amino a (B) TYPE: amino acid (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: S Met Gly His His His His His I  (2) INFORMATION FOR SEQ ID NO:18  (i) SEQUENCE CHARACTERISTIC (A) LENGTH: 20 base pa (B) TYPE: nucleic acid (C) STRANDEDNESS: sing (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: cDNA  (xi) SEQUENCE DESCRIPTION: S	Ile Val Gly Gly Glu Lys Ala Leu  1 5  Ser Leu Gln Ser Ser Ser His Phe 20  (2) INFORMATION FOR SEQ ID NO:17:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide  (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: SEQ ID  Met Gly His His His His His His His 1 5  (2) INFORMATION FOR SEQ ID NO:18: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: cDNA  (xi) SEQUENCE DESCRIPTION: SEQ I	Ile Val Gly Gly Glu Lys Ala Leu Ala  Ser Leu Gln Ser Ser Ser His Phe Ser  20 25  (2) INFORMATION FOR SEQ ID NO:17:  (i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 10 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide  (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: SEQ ID NO  Met Gly His His His His His His Glu 1 5  (2) INFORMATION FOR SEQ ID NO:18:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: cDNA  (xi) SEQUENCE DESCRIPTION: SEQ ID NO	Ser Leu Gln Ser Ser Ser His Phe Ser Gly 20 25  (2) INFORMATION FOR SEQ ID NO:17:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17: Met Gly His His His His His His Glu Phe 1 5 10  (2) INFORMATION FOR SEQ ID NO:18: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: cDNA  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	Ile Val Gly Gly Glu Lys Ala Leu Ala Gly Gln  Ser Leu Gln Ser Ser Ser His Phe Ser Gly Gly  20  (2) INFORMATION FOR SEQ ID NO:17:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide  (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:  Met Gly His His His His His His Glu Phe 1 5 10  (2) INFORMATION FOR SEQ ID NO:18:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: CDNA  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:	Ile Val Gly Gly Glu Lys Ala Leu Ala Gly Gln Ser 1 5 10  Ser Leu Gln Ser Ser Ser His Phe Ser Gly Gly Thr 20 25  (2) INFORMATION FOR SEQ ID NO:17:  (i) SEQUENCE CHARACTERISTICS:	The Val Gly Gly Glu Lys Ala Leu Ala Gly Gln Ser Pro  1 5 10  Ser Leu Gln Ser Ser Ser His Phe Ser Gly Gly Thr Ile 20 25  (2) INFORMATION FOR SEQ ID NO:17:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (C) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide  (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:  Met Gly His His His His His His Glu Phe 1 5 10  (2) INFORMATION FOR SEQ ID NO:18:  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(i) SeQUENCE CHARACTERISTICS: (A) LENGTH: 10 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: peptide  (v) FRAGMENT TYPE: internal  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:  Met Gly His His His His His His Glu Phe 1 5 10  (2) INFORMATION FOR SEQ ID NO:18:  (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear  (ii) MOLECULE TYPE: CDNA  (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

		Met L	ys Le	u Asn	Gln I	Lys 1	Asn i	Ala :		Ala 10	Val	Gly	Leu	Pro	A1a 15	Lys
5	(2)	INFORM	ATION	FOR S	SEQ I	D NO	:14:									
10	· <u></u>	(i) S	(A) I (B) 1	ICE CHA LENGTH: TYPE: &	: 7 a	mino aci	aci d	: ds								
		(ii) M	OLEC	JLE TY	PE: p	epti	de									
15		(v) F	TRAGMI	ENT TYI	PE: i	nter	nal									
		(xi) S	EQUE	NCE DES	SCRIP	TION	: SE	Q ID	NO:	14:						
20		Asn A	Asn G	ln Val	Val 5	Gly	Ile									
25	(2)	INFORM														
		(i) S	(A) :	NCE CH LENGTH TYPE:	: 31 amino	amin aci	o ac .d				:					
30		(ii) N		TOPOLO												
				ENT TY				al								
35		(xi) :	SEQUE	NCE DE	SCRII	PTION	1: SI	EQ II	ОИ	:15:						
÷		Ile '	Val G	ly Gly	Val	Lys	Ala	Lys	Ala	Gly 10	Asp	Ser	Pro	Tyr	Gln 15	Ile
40		Ser :	Leu G	ln Ser 20	Ser	Ser	His	Phe	Xaa 25	Gly	Gly	Ser	Ile	Leu 30	Asp	
45	(2)	INFOR	MATIC	N FOR	SEQ	ID N	0:16	:	/	Ź						
50		(i)	(A) (B)	ENCE CH LENGTH TYPE: TOPOLO	H: 30 amin	ami o ac	no a id	S: cids	;							
		(ii)	MOLEC	CULE TY	YPE:	pept	ide									
		(v)	FRAGN	MENT T	YPE:	N-te	rmin	al								

-41-

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Met Ile Cys Gly Gly Asp Val Ala Asn Gly Gly Val Asp Ser Glu Gln
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                         5
         Gly Asp
5
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10
               (A) LENGTH: 17 amino acids
               (B) TYPE: amino acid
               (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: peptide
15
         (v) FRAGMENT TYPE: internal
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20
         Met Ile Cys Gly Gly Asp Val Ala Asn Gly Gly Lys Asp Ser Cys Gln
                                             10
         Asp
25
     (2) INFORMATION FOR SEQ ID NO:12:
          (i) SEQUENCE CHARACTERISTICS:
30
               (A) LENGTH: 16 amino acids
               (B) TYPE: amino acid
               (D) TOPOLOGY: linear
         (ii) MOLECULE TYPE: peptide
35
         (v) FRAGMENT TYPE: internal
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:
         Met Thr Leu Asp Gln Thr Asn Ala Lys Pro Val Pro Leu Pro Thr Ser
40
                                              10
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45
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               (A) LENGTH: 16 amino acids
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50
         (ii) MOLECULE TYPE: peptide
          (v) FRAGMENT TYPE: internal
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:
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	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	
5	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
٠	TACATCCGAT CCTTTTGC	18
	TO T	
10	(Z) INFORMATION FOR SEQ ID NO:8:	
	(i) SEQUENCE CHARACTERISTICS:	٠,
	(A) LENGTH: 38 base pairs	
	<ul><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li></ul>	
15	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: cDNA	•
20	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
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25	(2) INFORMATION FOR SEQ ID NO:9:	
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	(A) LENGTH: 41 base pairs	
	(B) TYPE: nucleic acid	
30	(C) STRANDEDNESS: single	
	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: CDNA	
35	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
	GAATCAACAC CACCATTAGC AACATCACCA CCGCAAATCA T	41
40	(2) INFORMATION FOR SEQ ID NO:10:	
	AND ADDRESS OF THE PROPERTY OF	
	(i) SEQUENCE CHARACTERISTICS:  (A) LENGTH: 18 amino acids	
	(B) TYPE: amino acid	
45	(D) TOPOLOGY: linear	
	(ii) MOLECULE TYPE: peptide	
	(v) FRAGMENT TYPE: internal	
50	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	

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10		(ii) MOLECULE TYPE: cDNA	
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25		<ul><li>(B) TYPE: nucleic acid</li><li>(C) STRANDEDNESS: single</li><li>(D) TOPOLOGY: linear</li></ul>	
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40		(D) TOPOLOGY: linear	
		(ii) MOLECULE TYPE: cDNA	
45		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
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50	(2)	INFORMATION FOR SEQ ID NO:7:	
		<ul><li>(i) SEQUENCE CHARACTERISTICS:</li><li>(A) LENGTH: 18 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>	
55		(C) STRANDEDNESS: single	

		(i)	(A) (B)	LEN TYP	GTH: E: a	RACT 237 mino Y: 1	ami aci	no a d	: cids	ı							
5		(ii)	MOLE	CULE	TYP	E: p	epti	.de									
		(v)	FRAG	MENT	TYE	E: i	nter	mal									
10	-	(xi)															
		Ile 1	Val	Gly	Gly	Thr 5	Asp	Ala	Val	Leu	Gly 10	Glu	Phe	Pro	Tyr	Gln 15	Leu
15		Ser	Phe	Gln	Glu 20	Thr	Phe	Leu	Gly	Phe 25	Ser	Phe	His	Phe	Cys 30	Gly	Ala
••		Ser	Ile	Tyr 35	Asn	Glu	Asn	Tyr	Ala 40	Ile	Thr	Ala	Gly	His 45	Cys	Val	Tyr
20		Gly	Asp 50	Asp	туг	Glu	Asn	Pro 55	Ser	Gly	Leu	Gln	Ile 60	Val	Ala	Gly	Glu
25		Leu 65	Asp	Met	Ser	Val	Asn 70	Glu	Gly	Ser	Glu	Gln 75	Thr	Ile	Thr	Val	Ser 80
		Lys	Ile	Ile	Leu	His 85	Glu	Asn	Phe	Asp	Tyr 90	Asp	Leu	Leu	Asp	Asn 95	Asp
30		Ile	Ser	Leu	Leu 100	Lys	Leu	Ser	Gly	Ser 105		Thr	Phe	Asn	Asn 110	Asn	Val
25		Ala	Pro	Ile 115	Ala	Leu	Pro	Ala	Gln 120		His	Thr	Ala	Thr 125	Gly	Asn	Val
35		Ile	Val 130		Gly	Trp	Gly	Thr 135	Thr	Ser	Glu	Gly	Gly 140	Asn	Thr	Pro	Asp
40		Val 145		Gln	Lys	Val	Thr 150		Pro	Leu	Val	Ser 155	Asp	Ala	Glu	Cys	Arg 160
		Asp	Asp	Tyr	Gly	Ala 165		Glu	Ile	Phe	Asp 170	Ser	Met	Ile	Cys	Ala 175	Gly
45		Val	Pro	Glu	Gly 180	Gly	Lys	Asp	Ser	Cys	Ġln	Gly	Asp	Ser	Gly 190	Gly	Pro
		Leu	Ala	Ala 195		Asp	Thr	Gly	Ser 200		туг	Leu	Ala	Gly 205	Ile	val	Ser
50	•	Trp	Gly 210		Gly	/ Cys	: Ala	Arg 215		Gly	туг	Pro	Gly 220	Val	туг.	Thr	Glu
5.5		Val	. Ser	туг	r His	₅ Val	. Asp		ıle	e Lys	s Ala	Asn 235	Ala	. Val			

	Met -29	Ile	Ile	Tyr	Asn -25	Ile	Leu	Ile	Val	Leu -20	Leu	Leu	Ala	Ile	Asn -15	Thr
5 .	Leu	Ala	Asn	Pro -10	Ile	Leu	Pro	Ala	Ser -5	Pro	Asn	Ala	Thr	Ile 1	Val	Gly
10	Gly -	Glu 5	Lys	Ala	Leu	Ala	Gly 10	Glu	Cys	Pro	Tyr	Gln 15	Ile	Ser	Leu	Gln
10	Ser 20	Sèr	Ser	His	Phe	Cys 25	Gly	Gly	Thr	Ile	Leu 30	Asp	Glu	Tyr	Trp	Ile 35
15	Leu	Thr	Ala	Ala	His 40	Cys	Val	Ala	Gly	Gln 45	Thr	Ala	Ser	Lys	Leu 50	Ser
	Ile	Arg	Tyr	Asn 55	Ser	Leu	Lys	His	Ser 60	Leu	Gly	Gly	Glu	Lys 65	Ile	Ser
20	Val	Ala	Lys 70	Ile	Phe	Ala	His	<b>Gl</b> u 75	Lys	Tyr	Asp	Ser	Туг 80	Gln	Ile	Asp
25	Asn	Asp 85	Ile	Ala	Leu	Ile	Lys 90	Leu	Lys	Ser	Pro	Met 95	Lys	Leu	Asn	Gln
	Lys 100	Asn	Ala	Lys	Ala	Val 105	Gly	Leu	Pro	Ala	Lys 110	Gly	Ser	Asp	Val	Lys 115
30	Val	Gly	Asp	Gln	Val 120	Arg	Val	Ser	Gly	Trp 125	Gly	Tyr	Leu	Glu	Glu 130	Gly
	Ser	Tyr	ser	Leu 135	Pro	Ser	Glu -	Leu	Arg 140	Arg	Val	Asp	Ile	Ala 145	Val	Val
35	Ser	Arg	Lys 150	Glu	Cys	Asn	Glu	Leu 155	Tyr	Ser	Lys	Ala	Asn 160	Ala	Glu	Val
40	Thr	Asp 165	Asn	Met	Ile	Cys	Gly 170	Gly	Asp	Val	Ala	Asn 175	Gly	Gly	Lys	Asp
	Ser 180	_	Gln	Gly	Asp	Ser 185		Gly	Pro		Val 190		Val	Lys	Asn	Asn 195
45	Gln	Val	Val	Gly	Ile 200	Val	Ser	Trp	Gly	Tyr 205	Gly /	Cys	Ala	Arg	Lys 210	Gly
	Tyr	Pro	Gly	Val 215	Tyr	Thr	Arg	Val	Gly 220	Asn	Phe	Ile	Asp	Trp 225	Ile	Glu
50	Ser	Lys	Arg 230	Ser	Gln											

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	CAG	AAA	TAA	GCC	AAA	GCT	GTT	GGA	TTA	CCA	GCA	AAA	GGA	TCG	GAT	GTA		491
	Gln	Lys	Asn	Ala	Lys	Ala	Val	Gly	Leu	Pro	Ala	Lys	Gly	Ser	Asp	Val		
		100					105					110						
5	AAA	GTT	GGT	GAT	CAA	GTT	CGT	GTT	TCT	GGT	TGG	GGT	TAT	CTT	GAA	GAA		539
	Lys	Val	Gly	Asp	Gln		Arg	Val	Ser	Gly		GIÀ	Tyr	Leu	GIU	GIU		
	115					120					125					130		
							<b></b>	<i>(</i> (2) 2)	ምጥአ	מכת	ССТ	CTT	CAT	Σሞሞ	. GCT	GTT		, 587
	GGA	AGT	TAT	TCA	TIA	CCA	Cor	Clu	Lou	Ara	Ara	Val	Asp	Tle	Ala	Val		
10	GTA	ser	Tyr	ser		PIO	Ser	GIU	neu,	140	AL 9	***	ımp		145			-
					135	•												
•	CTA	TCA	ССТ	AAA	GAD	TGT	AAT	GAA	TTA	TAT	TCA	AAA	GCT	AAT	GCT	GAA		: 635
	Val	Ser	Δτα	LVS	Glu	Cvs	Asn	Glu	Leu	Tyr	Ser	Lys	Ala	Asn	Ala	Glu	-	
15	Val	501	9	150	<b>U</b>	-2-			155			_		160		. ,	11.	* .
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	GTT	ACT	GAT	AAT	ATG	ATT	TGT	GGT	GGT	GAT	GTT	GCA	AAT	GGT	GGT	AAA		683
	Val	Thr	Asp	Asn	Met	Ile	Cys	Gly	Gly	Asp	Val	Ala	Asn	Gly	Gly	Lys		
			165					170					175					
20																		777
	GAT	TCT	TGT	CAA	GGT	GAT	TCT	GGT	GGA	CCG	GTT	GTT	GAT	GTT	AAA	AAT		731
	Asp			Gln	Gly	Asp		GIA	GIÀ	Pro	vai	vai 190	Asp	vaı	гуя	Asn		
		180					185					190						
26	3 3 M	~ ~ ~ ~	C TTTT	CTT	CCT	א יוזיידי	CTT	י יירא	TGG	GGT	ТАТ	GGT	TGT	GCA	CGT	AAA		779
25	AAT	CAA	1751	Val	Glv	Tle	Val	Ser	Trp	Glv	Tvr	Glv	Cys	Ala	Arg	Lys		
	195		Val	VAI	Gry	200				1	205		•		•	210		
	GGT	TAT	CCA	GGT	GTT	TAT	ACA	CGT	GTT	GGT	AAT	TTT	ATC	GAT	TGG	ATT		827
30	Gly	Tyr	Pro	Gly	Val	Tyr	Thr	Arg	Val	Gly	Asn	Phe	Ile	Asp	Trp	Ile		
	_	_		_	215					220					225			
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				CGT				TTGA	AAT.	AAAC	AATT	TC G	ACAT	ATAA	A			8/5
	Glu	Ser	Lys	Arg		Gln												
35	*f* •			230	t .													
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		man n	eneren	ን ጥጥባ	יריי גיריי	יייי כי	CACC	מ מב) די	מ מ	'AAAC	'AATC	. ATG	ATTG	TAT	AGGG	ATTT	3T	995
40	TTA	CAT	111	MIII	IMIA	.11 C	.GAGC											
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	. 131																	
	AAA	.G					•											1059
		-																

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- (2) INFORMATION FOR SEQ ID NO:2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 261 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
    - (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

5			(E	LE 3) TY C) ST () TO	PE:	nucl EDNE	eic SS:	acid sing	ļ	s								-
٠		(ii)	MOI	ECUL	E TY	PE:	CDNA											
10	-	_(ix)		ATURE A) NA B) LC	ME/K			848										
15		(ix)	-	ATURE A) NA B) LC	ME/K		_	_			,							
		(xi)	SEÇ	QUENC	E DE	SCRI	PTIC	N: S	SEQ I	D NO	):1:							
20	CTT	rttt:	rtt 1	CTTTI	TGAF	A AC	ATTO	AAA	ACA	TTGF	AAGC	AACI	GTGT	GT A	CAAC	CAGA	A	60
25	. 1		ATC A		yr I					al I					le A			107
25			GCT Ala															155
30			GAA Glu 5															203
35			AGT Ser														·	251
40			ACA Thr													CTT Leu 50	:	299
			CGT Arg															347
45	TCT Ser	GTT Val	GCT Ala	AAA Lys 70	ATT Ile	TTT Phe	GCA Ala	CAT His	GAA Glu 75	AAA Lys	TAT Tyr	GAT Asp	AGT Ser	TAT Tyr 80	CAA Gln	ATT		395
50			GAT Asp 85	Ile					Leu									443

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### SEQUENCE LISTING

	(1) GENERAL INFORMATION:	
5	(1)	
•	(1) APPLICANT: (A) NAME: IMMULOGIC PHARMACEUTICAL CORPORATION	
	(B) STREET: 610 LINCOLN STREET	
	(C) CITY: WALTHAM	
10	(D) STATE: MASSACHUSETTS	
10	(E) COUNTRY: USA	
	(F) POSTAL CODE (ZIP): 02154	
	(G) TELEPHONE: (617) 466-6000	
	(H) TELEFAX: (617) 466-6010	
15	(ii) TITLE OF INVENTION: NUCLEIC ACIDS ENCODING A HOUSE DUST MI ALLERGEN, DER P III, AND USES THEREFO	T R
	(iii) NUMBER OF SEQUENCES: 18	
20		
	(iv) CORRESPONDENCE ADDRESS:	
	(A) ADDRESSEE: LAHIVE & COCKFIELD (B) STREET: 60 STATE STREET, SUITE 510	
	(B) STREET: 60 STATE STREET, SOTTE SEC	
	(D) STATE: MA	
25	(E) COUNTRY: USA	
	(F) ZIP: 02109	
	(v) COMPUTER READABLE FORM:	
30	(A) MEDIUM TYPE: Floppy disk	
	(B) COMPUTER: IBM PC compatible	
	(C) OPERATING SYSTEM: PC-DOS/MS-DOS	
	(D) SOFTWARE: ASCII TEXT	
35	(vi) CURRENT APPLICATION DATA:	
<i>33</i> , .	(A) APPLICATION NUMBER:	
	(B) FILING DATE:	
	(C) CLASSIFICATION:	
	TO THE TOTAL PROPERTY OF THE PARTY.	
40	<pre>(vii) PRIOR APPLICATION DATA:    (A) APPLICATION NUMBER: US 08/163,919</pre>	
	(B) FILING DATE: 08 DEC 1993	
	(B) FIBING DATE: OF HE	
	(viii) ATTORNEY/AGENT INFORMATION:	
45	(A) NAME: MANDRAGOURAS, AMY E.	
	(B) REGISTRATION NUMBER: 36,207/	
	(C) REFERENCE/DOCKET NUMBER: 072.0US(IMI-041PC)	
	(ix) TELECOMMUNICATION INFORMATION:	
50	(A) TELEPHONE: (617) 227-7400	
20	(B) TELEFAX: (617) 227-5941	

(2) INFORMATION FOR SEQ ID NO:1:

molecular weight of approximately 32 kDa is slightly higher than that predicted from the primary structure (Figure 1A and 1B, and above). In Figure 3, increasing concentrations of Der p III preparation were examined to verify the purity (lane 1, 4.3  $\mu$ g; lane 2, 8.7  $\mu$ g; lane 3, 13.8  $\mu$ g; lane 4, 17.4  $\mu$ g; markers are indicated by M).

**Equivalents** 

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Those skilled in the art will recognize or be able to ascertain using no more than routine experimentation, numerous equivalents to the specific embodiments described herewith. Such equivalents are considered to be within the scope of this invention and are encompassed by the following claims.

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(MacDonald et al. J Biol Chem (1982) 257:9724-9732). Stewart et al. (1992) proposed nine major isoforms of Der p III existed, with pI's ranging from 4 to >8.

# EXAMPLE 3: Expression and Purification of Recombinant Der p III

A complementary DNA insert encoding Der p III (Figure 1A and 1B) was digested 5 with EcoRI and inserted into pUC19. To remove the non-translated 5' sequence as well as part of the putative hydrophobic leader, the cDNA was cleaved with MscI and an EcoRI linker was added. Subsequently, the truncated Der p III fragment was EcoRI digested and subcloned into the expression vector pET11d (Studier et al., Gene Expression Technology: Methods in Enzymology 185, Academic Press, San Diego, California (1990) 60-89). The 10: sequence and reading frame were verified by DNA sequencing. The Der p III coding sequence encompassed residue 13 to the stop codon (see Figure 1A and 1B). The expression vector pET11d Der p III was transformed into the E. coli host strain BL21(DE3) and selected on plates containing 150 µg/ml ampicillin (Studier et al., Gene Expression Technology: Methods in Enzymology 185, Academic Press, San Diego, California (1990) 60-89). A 15 single transformant colony was grown up in 2 ml volume of BHIB medium containing 150 μg/ml ampicillin at 37°C, for approximately 6 hr. Ten milliliters of this culture was spread onto a selection plate and grown overnight at 37°C. The bacterial lawn was recovered in 2 ml media and added to 500 ml of BHIB medium (150 μg/ml ampicillin) and grown at 37°C to A600=1.0. Recombinant expression was induced by the addition of IPTG to a final 20 concentration of 1 mM. After 2 hr growth, cells were harvested, lysed and the proteins solubilized in 6 M guanidine HCl buffer containing 100 mM 2-ME as previously described for the ragweed recombinant allergens, Amb a I.1 and Amb a II (Rogers, et al. (1991) J. Immunol. 147:2547-2552).

The guanidine HCl lysate containing *Der p* III was subjected to Ni<sup>2+</sup> metal- ion affinity chromatography under denaturing conditions in 8 M urea (Hochuli, E. et al., (1988) Bio/Technology 6:1321-1325). After elution from the Ni<sup>2+</sup> chelating support, QIAGEN NTA-Agarose (Diagen GmH, Dusseldorf, Germany), the recombinant *Der p* III protein preparations were subjected to SDS-PAGE analysis (Figure 3).

Recombinant *Der p* III had an expected molecular weight of approximately 27 kDa as predicted for the 252 amino acids plus amino acids used as a purification tag (MGHHHHHHEF (SEQ ID NO:17)). The pET11d expression system and the Ni<sup>2+</sup> methylion-affinity chromatography method generated approximately 12 mg recombinant *Der p* III per liter of growth medium (as assessed by A<sub>280</sub> measurement). This expression and purification scheme yielded a protein preparation with purity exceeding 90% as assessed by SDS-PAGE visualized by Coomassie Blue staining. (Figure 3). The observed apparent

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1971, pages 249-279). The glycine residues at 206 and 211 in combination with the aspartic acid at 179 are responsible for accommodating the bulky positively charged side chains of lysine and arginine, the substrate residues cleaved by trypsins. Amino acid residues 179-185 and 202-206 (Figure 1A and 1B) form the S1 binding pockets which are involved in the binding of the substrate to the trypsin molecule. The areas immediately surrounding all of these important residues are the most conserved regions of the *Der p* III protein. All these results are consistent with the fact that among trypsin proteins from different species structural variations occur in those regions of the molecule which are not important for catalytic activity (Vithayathil et al., Arch Biochem Biophys (1961) 92:532-540).

Most trypsin proteins generally have a common pairing of cysteine residues to form six disulphide bridges, with two of these bridges, Cys 15-145 and Cys 117-218 being unique to this group of proteins. Der p III like both crayfish (Titani et al., Biochemistry (1983) 22:1459-1465) and S. griseus are missing these two unique links, but more importantly Der p III is unique as it contains an extra unpaired cysteine residue. This residue may be the equivalent to Cys 15 in the bovine trypsin which unlike, Cys 145, 117 and 218 was not eliminated during evolutionary divergence of the mite trypsin.

The isolation of the gene encoding the nucleotide sequence for the group III allergen, Der p III is the first primary sequence determination for this group of Dermatophagoides allergens. Comparison of the sequence with other vertebrate and invertebrate serine proteases in view of substrate binding experiments (Stewart et al., Immunology (1992) 75: 29-35; Ando et al., Arerugi (1992) 41:704-707), indicates that the group III allergens are trypsin proteins. Trypsin proteins are secreted as pre-pro zymogens by the pancreatic acinar cells of all vertebrate and invertebrate species. Invertebrate trypsins have been reported to have a molecular weight ranging from 20 to 30kDa (Graf et al., Insect Biochem (1985) 15:611-618). The Der p III P3WS1 trypsinogen has a calculated molecular weight of 28kDa and the corresponding mature protein a molecular weight of 24,985kDa while the native protein purified, existed as duplicate of 28 and 30kDa as estimated by SDS-PAGE. There have been other reports of multiple protein bands being isolated for both Derf III (Thomas et al., Exp Appl Acarol (1992) 16:153-164) and Der p III (Stewart et al., Immunology (1992) 75: 29-35). These results are not unusual as trypsin proteins are known to exist in several isoforms. Gendry and Launay (1988) isolated duplicate protein bands of an anionic trypsinlike protein from rat pancreas. They demonstrated that the higher 32kDa band represented the inactive trypsin and the lower 30kDa band represented the active form of the trypsin which could autocatalyze the cleavage of the 32kDa band. It is therefore possible that the 30kDa band represents the inactive trypsinogen and the 28kDa band the active trypsin. The number of isozymes for invertebrate trypsins has been reported to range from 1-12

for both *Drosophila melangastor* (Davis et al., Nucleic Acid Research (1988) 13:6605-6615) and S. griseus (Kim et al., Biochem Biophys Res Commun (1991) 181:707-713). It is possible therefore that the highly conserved clusters of hydrophobic residues are not a characteristic of all trypsin proteins but rather that of mammalian trypsins.

Downstream from the signal peptide is the 11 residue pre-activation peptide from -:: 5 amino acid residues -11 to -1. Comparison of this pro region with other pro regions from arrypsins of various species indicates the Der p III pro peptide is unusual. Both vertebrate and invertebrate trypsins have been shown to have an octapeptide and or a hexapeptide pro region with four contiguous aspartyl residues followed by a lysine residue at the -1 position (Le Huerou et al., Eur J Biochem (1990) 193:767-73). It is the carboxyl region of this lysine residue which is cleaved during the activation of the trypsinogen to trypsin. The Der p III lacks this polyaspartyl-lysine sequence. The trypsinogen therefore can not be activated by the usual mechanism. There are a few other trypsins which are similar to Der p III in this respect, D. melangastor (Davis et al., Nucleic Acid Research (1988) 13:6605-6615), S. griseus (Kim et al., Biochem Biophys Res Commun (1991) 181:707-713) and most 15 interestingly, a 32kDa anionic trypsin from the rat pancreas (Gendry and Launay, Biochemica et Biophysica Acta (1988) 955:243-249). This group showed that enterokinase, an enzyme which has a highly specific affinity for the polyaspartyl-lysine residues in trypsin proteins, had no effect on the trypsin that they isolated. They suggested therefore that a different mechanism was being used for the activation process. The absence of any 20 homology of the Der p III proenzyme region with any of the known sequences may imply a unique activation mechanism.

A search of the protein database has confirmed that Der p III P3WS1 is homologous .154 Tab. to both vertebrate and invertebrate trypsins. The Der p III allergen is not a mite chymotrypsin (Gendry and Launay, Biochemica et Biophysica Acta (1988) 955:243-249) as comparison of the N-terminal protein sequences showed that they differ by as much as 50%. Figure 2A and 2B show a comparison of the amino acid sequences of Der p III P3WS1 and crayfish trypsins (SEQ ID NOS: 2 and 3). The crayfish has 44% homology with Der p III. There are seven cysteine residues, six of these residues are known to form disulphide bridges, 54-70, 181-198 and 210-236 (Figure 2A and 2B.) (Hartley, Phil Trans Roy Soc Lond B 30 (1970) 257:77-87). Most importantly, the Der p III protein contains the highly conserved residues involved in the catalytic activity and substrate specificity of the trypsin proteins. The conserved His 40 and Ser 185 (Figure 1A and 1B) represent the charge relay system which comprise the active catalytic site of the trypsin enzyme. The positively charged aspartic acid residue at 179, represents the trypsin specificity site. Most serine proteases 35 have a neutral amino acid at this residue (Keil, The Enzymes. Academic Press, New York,

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These substitutions may be due to the existence of Der p III in different isoforms. Analysis of the deduced protein sequence for Der p III P3WS1 confirmed that the position of the Der f III CNBr peptide 1 (MI(C)GGDVANGGVDS(E)QGD) (SEQ ID NO:10) was correctly - predicted to be from amino acid residues 177-183 and has 88% identity with the recombinant sequence in this region, MICGGDVANGGKDSCQD (SEQ ID NO:11). Interestingly, there are two identical non-conservative changes that exist when the native Der f III peptide sequence is aligned with the P3WS1 clone sequence. There is a change from a non-polar hydrophobic valine at 178 (according to numbering in Figure 1A and 1B) in Der f III to a positively charged lysine residue in the P3WS1 clone sequences. The other change is a glutamic acid residue at 181 in the Der f III substituted for a cysteine residue in the P3WS1 clone sequence. The cysteine forms one of the disulfide bridges in both these and most other trypsin proteins. The predicted alignment of the second Der f III peptide, (M)TLDQTNA(K)PVPL(P)(T)(S) (SEQ ID NO:12), produced by the CNBr digestion was from amino acid residues 95-110 (MKLNQKNAKAVGLPAK) (SEQ ID NO:13). There is 63% homology with the P3WS1 clone in this region. These differences between the Der pIII and Der f III protein sequences are consistent with the 20% sequence variation found between other homologous allergens from this species such as the Der p I and Der f I; and Der p II and Der f II.

The complete sequence for the Der p III P3WS1 clone encodes a protein of 319 amino acids. Der p III, like all known trypsins, is synthesized as a pre-pro zymogen. The cleavage site of the signal peptide is postulated to be between amino acid residues -12 and -11 as the residues around this site conform to the amino acid constraints outlined by von Heijne, (von Heijne, Eur J Biochem (1983) 133:17-21). Von Heijne proposed that small neutral amino acids are strongly preferred at positions -1 and -3 from the cleavage site, such as the alanine and tyrosine seen in the Der p III gene. Proline residues are never found in positions +1 to -3 thus excluding any other sites. Most mammalian trypsin proteins have been reported to have pre or signal peptides of 15 or 16 amino acids. The Der p III peptide is 18 amino acids in length. While the difference in length is relatively small, it does seem that the length of the signal peptide may be attributable to the phylogenetic diversity of the species from which the trypsin originated. This is most apparent in Streptomyces griseus with a 32 amino acid pre peptide (Kim et al., Biochem Biophys Res Commun (1991) 181:707-713). It has also been reported that for mammalian trypsins the signal peptide contains two specific clusters of two and then four hydrophobic residues (MacDonald et al., J Biol Chem (1982) 257:9724-9732; Le Huerou et al., Eur J Biochem (1990) 193:767-73). There is an abundance of hydrophobic residues within this region of the Der p III gene but

There is an abundance of hydrophobic residues within this region of the *Der p* III gene but they are not arranged into highly conserved clusters. The same is true of the signal peptides

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sequence. The Der p III N-terminal sequence was extended to IVGGEKALAGQSPYQISLQSSSHFSGGTIL (SEQ ID NO:16). The Der f III N-terminal sequence was extended to IVGGVKAKAGDSPYQISLQSSSHFXGGSILD (SEQ ID NO:15). Comparison of the sequence data published by Stewart et al. (Immunology (1992) 75: 29-35) and Heymann et al., (J. Allergy. Clin. Immunol (1989) 83: 1055-1067) for the group III allergens with the data herein indicated several errors in the published sequence. In particular, a non-conservative change at residue 8 from a positively charged lysine to a non-polar hydrophobic leucine was found (Der f III, Heymann et al., J Allergy Clin Immunol (1989) 83: 1055-1067). Two internal peptides of Der f III were isolated by HPEC after CNBr digestion of the natural protein, MICGGDVANGGVDSEQGD (SEQ ID NO:10) and MTLDQTNAKPVPLPTS (SEQ ID NO:12).

## EXAMPLE 2: Sequence of recombinant Der p III

The protein sequence information obtained as described in Example 1 was used in conjunction with the preferred codon usage data (Table 1) to construct two oligonucleotides probes. These oligonucleotides were designed to hybridize to nucleotide residues 159-196 and to residues 688-648 of the Der p III clone (Figure 2A and 2B). Only clones which hybridized strongly with both probes were isolated from the  $\lambda$  gt10 cDNA library. The resulting nucleotide sequence for the P3WS1 clone and the deduced amino acid sequence is shown in Figure 1A and 1B. The complete nucleotide sequence was 1059 bp in length. This includes a 5' non-coding region of 62bp, a 211bp 3' untranslated region and an open reading frame of 786 bp with a stop codon (TGA) at nucleotide residues 846-848. There is no poly A tail but there does appear to be a polyadenylation signal (AATAAA). The open reading frame encodes a protein which includes a 29 amino acid pre-pro region and starting at the Nterminal isoleucine, a mature protein of 232 amino acid residues with a calculated molecular weight of 24,985 and pI of 8.5. The methionine residue (ATG) at amino acid position -29 is the most likely translation initiation site. Selection for this initiation site was based in part on the presence of a sequence following the methionine which encodes a classical signal peptide of 18 amino acid residues with predominantly hydrophobic residues present. In addition, the immediate sequence prior to this ATG codon, 5'GAAAGATG3', conforms loosely to the Kozak consensus sequence (CCACCATG) for the eucaryotic translation initiation site with the crucial purine (most often A) at the -3 position (Kozak, Nucleic Acid Research (1984) 12:857-872).

The protein sequencing data of *Der p* III described in Example 1 differs from the amino acid sequence deduced from the cDNA of the P3WS1 clone as a result of the substitutions of Glu/Gln at residue 11 and a possible Cys/Ser substitution at residue 17.

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#### Isolation of DNA from λ gt10 Der p III Clones

Phage DNA from the *Der p* III clones was prepared using a polyethylene glycol precipitation procedure (Chua <u>et al.</u>, <u>J Exp Med</u> (1988) <u>167</u>:175-182).

### Subcloning and DNA Sequencing

Purified *Der p* III phage DNA was digested with EcoRI restriction enzyme (Toyobo) and the released fragment ligated to EcoRI restriction enzyme digested M13mp19 sequencing vector (Messing, Meth. Enzymol. (1983) 101: 20-78). The recombinant DNA was transformed using *E. coli* TG-1 and sequencing performed using the dideoxynucleotide chain termination method and Sequenase (U.S. Biochemicals) (Sanger et al., Proc Natl Acad Sci USA (1977) 74:5463-5467). The oligonucleotide primers used for sequencing included the universal primers, a 17-mer sequencing primer (-40)5' CAGCACTGACCCTTTTG3' (SEQ ID NO:4) and a 16-mer reverse sequencing primer (-24)5' AACAGCTATGACCATG3' (SEQ ID NO:5), the two internal primers used for the library screening (SEQ ID NOS: 8 and 9) and two internal primers P3forward5(P3F5) 5'AAAGCTGTTGGATTACCA3' (SEQ ID NO:6) and P3reverse5(P3R5) 5'TACATCCGATCCTTTTGC3' (SEQ ID NO:7) and P3F4
5'GCGGATCCATTGTTGGTGGT 3' (SEQ ID NO:18). The two internal probes were designed to correspond to nucleotide residues 456-473 (P3F5) and 491-474 (P3R5) (Figure 1A and 1B). The primers were used to sequence the isolated clone in both orientations.

#### DNA and Protein Sequence Analysis

Sequence analysis was performed using the MAC VECTOR software (IBI, New Haven, U.S.A.). Computation for sequence homology with other proteins was performed at the NCBI using the BLAST network service. The version of BLAST used was BLAST 1.3.10MP (7-Jul-93).

#### **EXAMPLE 1: Sequence of the native group III allergens**

The protein isolation procedure produced a *Der p* III sample which ran as a doublet with molecular weights 30 and 28k when analyzed by SDS-PAGE. Both bands reacted with polyclonal mouse anti-*Der f* III in accord with the interspecies cross-reactivity previously reported (Thomas et al., Exp. Appl. Acarol. (1992) 16:153-164). *Der f* III isolated from a 5A12 monoclonal antibody (Heymann et al. J. Allergy Clin. Immun. (1989):1055-1067) column exhibited similar characteristics. Using the o-pthalaldehyde to eliminate the contaminating protein sequences, the N-terminal sequencing of both native *Der p* III and *Der f* III corrected and extended the known *Der p* III sequence and extended the *Der f* III

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## Construction of the D. pteronyssinus λ gt10 cDNA Library

Polyadenylated mRNA (10μg) was used to synthesize cDNA by the RNase H method (Gubler et al. Gene (1983) 25:263-269) using a kit (Amersham International, Aylesbury, U.K.). After the addition of EcoRI restriction enzyme linkers (New England Biolabs, Beverly, U.S.A.), the cDNA was ligated to alkaline-phosphatase treated lambda gt10 arms (Promega Biotec, Madison, Wisc.). The recombinant phage DNA was packaged and plated wout in E. coli JP777 to produce a library of 5x10<sup>5</sup> recombinants.

### Kinase End-labeling of Oligonucleotides

Oligonucleotides were synthesized using an Applied Biosystem PCR mate (Applied Biosystems, Foster City, CA). Twenty picomole of oligonucleotide DNA was end labeled with (γ -32P) ATP using T4 polynucleotide kinase (Promega Corp., Madison, Wisc.) (Maniatis, T., Fritsch, E. F. and Sambrook, J. Molecular Cloning: A Laboratory Manual. 2nd Ed. Cold Spring Harbor Laboratory Press, 1989). Labeled oligonucleotide was purified by 15% polyacrylamide gel electrophoresis and subsequently eluted into sterile distilled water.

## Isolation of Der p III cDNA Clones from the λ gt10 cDNA Library

Screening of the library was performed with two probes designed using both N-terminal and internal protein sequence data obtained as described above for the group III allergens. The first, P3forward3 (P3F3), was a 38mer in length with the following nucleotide sequence, 5'TCAGAAAAAGCTTTGGCTGGTGAATCACCATATCAAAT 3' (SEQ ID NO:8). The second probe, P3reverse4(P3R4), a 41mer with the following nucleotide sequence, 5'GAATCAACACCACCATTAGCAACATCACCACCGCAAATCAT3' (SEQ ID NO:9). The library was plated at 25,000 pfu per 150mm petri dish and the phage were lifted onto nitrocellulose (Schleicher & Schuell, Dassel, Germany). Duplicate filter lifts from each plate were denatured and baked for hybridization with a different one of the two probes (Maniatis, T., Fritsch, E. F. and Sambrook, J. Molecular Cloning: A Laboratory Manual. 2nd Ed. Cold Spring Harbor Laboratory Press, 1989). Hybridizations were performed in hybridization mix (6x sodium chloride/sodium citrate pH 7.0 (SSC), 0.1% Denhardts, 100µ g/ml denatured herring sperm DNA) at 42°C with 10<sup>6</sup>cpm/ml labeled probe overnight. Filters were washed three times at 42°C for 20 min and then at 50°C for 10 min, in 400 ml wash solution containing 6x SSC, 0.1% Triton X-100.

Table 1. Preferred codon usage table for Dermatophagoides allergens.

Ala       A       GCT       45.4       Pro       P       CCA       87.2         GCC       33       CCC       9.6         GCA       21.6       CCC       9.6         GCG       0       CCT       0         Cys       C       TGT       55.2       Gln       Q       CAA       94.4         Cys       C       TGC       44.8       Gln       Q       CAA       94.4       CAG       5.6         Asp       D       GAT       92.3       Arg       R       CGA       36.5       CGT       41.6       CGC       16.5       CGT       41.6       CGC       16.5       CGC       16.5       AGA       5.4       CGC       16.5       AGA       5.4       CGC       16.5       AGA       5.4       CGC       16.5       AGA       5.4       CGG       0       AGA       5.4       CGG       0       AGA       5.4       CGG       0       AGA       5.4       CGG       0       AGA       10.2       CGG       0       AGA       11.2       AGA       11.2       AGA       11.2       AGA       11.2       AGA       11.2       AGA       13.5       ACC	Amin	o acid	l Codon	%		Amino	n acid	Codon	%
GCC GCA 21.6 CCG 3.2 CCT 0  Cys C TGT 55.2 Gln Q CAA 94.4 CAG 5.6  Asp D GAT 92.3 Arg R CGA 36.5 CGT 41.6 CGC 16.5  Glu E GAA 100 AGA 5.4 CGG 0  GAG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 TCT 18.4 AGT 11.2 GGA 24.5 GGC 10.3 AGC 10.2 GGG 0  His H CAT 78.8 Thr T ACT 39.2 ACA 35 ACC 18.9 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 AGG 1 CGT 33.8  Lys K AAA 99 AGG 1  Leu L TTG 64.5 Trp W TGG 100  CCG 3.2  GGA 24.5 GTA 12.3  GTA 12.3  GTG 2.7  Leu L TTG 64.5 Trp W TGG 100									
GCA GCG 0 CCT 0  Cys C TGT 55.2 Gin Q CAA 94.4  Asp D GAT 92.3 Arg R CGA 36.5  GAC 7.7 CGT 41.6  Glu E GAA 100 AGA 5.4  GAG 0  Phe F TTC 82.5  TTT 17.5 Ser S TCA 42.5  GGG 0  GGG 10.3  GGG 0  His H CAT 78.8 Thr T ACT 39.2  GGG 10.2  GGG 10.2  GGG 6.5  His H CAT 78.8 Thr T ACT 39.2  CAC 21.2  ACA 35  ACC 18.9  ACC 18.9  Lys K AAA 99  AAG 1  Lys K AAA 99  AAG 1  Leu L TTG 64.5 Trp W TGG 100  CCC 3.3.2  CCA 94.4  GGA 94.4  CAG 3.2  CCA 94.4  CCG 10.2  CCG 16.5  CGT 41.6  CGG 0  AGG 0  AGG 10.2  TCC 6.5  TTT T ACT 39.2  ACC 18.9  GTA 12.3  GTG 2.7	Ala	A					•		
GCG 0 CCT 0  Cys C TGT 55.2 Gin Q CAA 94.4 TGC 44.8 CAG 5.6  Asp D GAT 92.3 Arg R CGA 36.5 GAC 7.7 CGT 41.6 CGC 16.5  Glu E GAA 100 AGA 5.4 CAG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 GGA 24.5 GGC 10.3 GGC 10.3 GGG 0 TCC 6.5  His H CAT 78.8 Thr T ACT 39.2 GGG 10.2 GGG 6.9  ATC 48.5 ATA 1.5 Val V GTT 51.2 GTA 12.3 GTG 33.8 Lys K AAA 99 AAG 1  Leu L TTG 64.5 Trp W TGG 100									
Cys         C         TGT TGC         55.2 t44.8         Gln Q CAA CAG         94.4 CAG         5.6           Asp D GAT GAC 7.7         92.3 Arg R CGA 36.5 CGT 41.6 CGC 16.5         CGT 41.6 CGC 16.5         CGC 16.5         CGC 16.5         CGG 0 CGG 11.2 CGG									
TGC 44.8 CAG 5.6  Asp D GAT 92.3 Arg R CGA 36.5 GAC 7.7 CGT 41.6 CGC 16.5 Glu E GAA 100 AGA 5.4 GAG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 GGA 24.5 GGA 24.5 GGG 10.3 GGG 0 TCC 6.5  His H CAT 78.8 Thr T ACT 39.2 CAC 21.2 ACA 35 ACC 18.9 Ile I ATT 50 ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 AAG 1  Leu L TTG 64.5 Trp W TGG 100  TCG 100  TCG 11.2 TCG 33.8 Thr 50 ACG 6.9 ATC 48.5 ATA 1.5 Trp W TGG 100			GCG					CC1	. 0
Asp D GAT 92.3 Arg R CGA 36.5 GAC 7.7 CGT 41.6 CGC 16.5 GIu E GAA 100 AGG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 TCT 18.4 GGA 24.5 GGC 10.3 AGC 10.2 GGG 0 TCC 6.5 GGG 11.2 AGC 12.2 AGC 12.2 AGC 13.9 ACC 18.9 IIe I ATT 50 ACC 18.9 IIe I ATT 50 ACC 18.9 ACC 18.9 Lys K AAA 99 GTA 12.3 AGC 1 CGG 2.7 Leu L TTG 64.5 Trp W TGG 100 Low TCC 10.0 CGG 10	Cys	C	TGT	55.2		Gin	Q	ÇAA	94.4
GAC 7.7 CGT 41.6 CGC 16.5  Glu E GAA 100 AGA 5.4 CGG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 GGA 24.5 GGC 10.3 GGG 0 TCC 6.5  His H CAT 78.8 Thr T ACT 39.2 CAC 21.2 ACA 35 ACC 18.9 Ile I ATT 50 ACG 6.9 ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 Lys K AAA 99 Leu L TTG 64.5 Trp W TGG 100  Leu L TTG 64.5 Trp W TGG 100  CGC 16.5  41.6 CGC 16.5  AGA 5.4 CGG 0 AGA 5.4 CCG 0 AGA 42.5 TCT 18.4 A2.5 TCT 39.2 ACC 18.9 ACC 18.9 ACC 33.8 ACC 33.8 ACC 33.8 ACC 12.3 ACC 33.8 ACC 33.		•	TGC	44.8				CAG	5.6
GAC 7.7 CGT 41.6 CGC 16.5  Glu E GAA 100 AGA 5.4 CGG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 GGA 24.5 GGC 10.3 GGG 0 TCC 6.5  His H CAT 78.8 Thr T ACT 39.2 CAC 21.2 ACA 35 ACC 18.9 Ile I ATT 50 ACG 6.9 ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 Lys K AAA 99 Leu L TTG 64.5 Trp W TGG 100  Leu L TTG 64.5 Trp W TGG 100  CGC 16.5  41.6 CGC 16.5  AGA 5.4 CGG 0 AGA 5.4 CCG 0 AGA 42.5 TCT 18.4 A2.5 TCT 39.2 ACC 18.9 ACC 18.9 ACC 33.8 ACC 33.8 ACC 33.8 ACC 12.3 ACC 33.8 ACC 33.	Asp	D	GAT	92.3		Arg	R	CGA	36.5
Glu E GAA 100 AGA 5.4  GAG 0 CGG 0 AGG 0  Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 TCT 18.4  Gly G GGT 65.2 AGT 11.2 GGA 24.5 TCG 11.2 GGA 10.3 AGC 10.2 GGG 0  His H CAT 78.8 Thr T ACT 39.2 CAC 21.2 ACA 35  Ile I ATT 50 ACA 35  ATA 1.5 Val V GTT 51.2 GTC 33.8  Lys K AAA 99 GTA 12.3 AAG 1  Leu L TTG 64.5 Trp W TGG 100  Leu L TTG 64.5 Trp W TGG 100  TTC 100  TTA 18.5		_			•	J			
Glu       E       GAA       100       AGA       5.4         GAG       0       CGG       0         Phe       F       TTC       82.5       TCA       42.5         TCT       118.4         Gly       G       GGT       65.2       AGT       11.2         GGA       24.5       TCG       11.2         GGC       10.3       AGC       10.2         GGG       0       TCC       6.5         His       H       CAT       78.8       Thr       T       ACT       39.2         ACA       35       ACC       18.9         Ile       I       ATT       50       ACG       6.9         ATA       1.5       Val       V       GTT       51.2         GTC       33.8         Lys       K       AAA       99       GTA       12.3         AAG       1       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100	:			• • • • • • • • • • • • • • • • • • • •	•				
GAG	Gl.,	E	GAA	100					
Phe F TTC 82.5 TTT 17.5 Ser S TCA 42.5 TCT 18.4 AGT 11.2 AGT 11.2 AGC 10.2 AGC 10.2 AGC 10.2 AGC 10.2 ACC 18.9 ACC 18.9 ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 GTA 12.3 AAG 1 CTC 10.0 TTA 18.5	Giu	E		•					
Phe         F         TTC         82.5           TTT         17.5         Ser         S         TCA         42.5           TCT         18.4           Gly         GGT         65.2         AGT         11.2           GGA         24.5         TCG         11.2           GGC         10.3         AGC         10.2           GGG         0         TCC         6.5           His         H         CAT         78.8         Thr         T         ACT         39.2           CAC         21.2         ACA         35         ACC         18.9           Ile         I         ATT         50         ACG         6.9           ATA         1.5         Val         V         GTT         51.2           GTC         33.8         GTA         12.3         GTA         12.3           Lys         K         AAA         99         GTA         12.3           AAG         1         GTG         2.7    Leu  L  TTG  64.5  TTA  TTA  TTA  TTD  W  TGG  100			GAG	·					
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Gly       G       GGT       65.2       AGT       11.2         GGA       24.5       TCG       11.2         GGC       10.3       AGC       10.2         GGG       0       TCC       6.5         His       H       CAT       78.8       Thr       T       ACT       39.2         CAC       21.2       ACA       35       ACC       18.9         Ile       I       ATT       50       ACG       6.9         ATC       48.5       ATA       1.5       Val       V       GTT       51.2         GTC       33.8       GTA       12.3       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100         TTA       18.5       TTP       W       TGG       100									18.4
GGA 24.5 TCG 11.2 GGC 10.3 AGC 10.2 GGG 0 0 TCC 6.5  His H CAT 78.8 Thr T ACT 39.2 ACA 35 ACC 18.9 ACC 18.9 ACG 6.9 ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 GTA 12.3 AAG 1 GTG 2.7  Leu L TTG 64.5 Trp W TGG 100 TTA 18.5	Glv	G	GGT	65.2					
GGC 10.3 AGC 10.2 TCC 6.5  His H CAT 78.8 Thr T ACT 39.2 ACA 35 ACC 18.9  Ile I ATT 50 ACG 6.9  ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8  Lys K AAA 99 GTA 12.3 AAG 1 GTG 2.7  Leu L TTG 64.5 Trp W TGG 100	,	_							
His       H       CAT CAC       78.8 CAC       Thr       T       ACT ACA ACA ACA ACC ACC ACC ACC ACC ACC									
His H CAT 78.8 Thr T ACT 39.2 CAC 21.2 ACA 35 ACC 18.9 Ile I ATT 50 ACG 6.9 ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Lys K AAA 99 GTA 12.3 AAG 1 GTG 2.7 Leu L TTG 64.5 Trp W TGG 100									
Ile       I       ATT       50       ACG       18.9         ATC       48.5       ACG       6.9         ATA       1.5       Val       V       GTT       51.2         GTC       33.8       GTA       12.3         AAG       1       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100         TTA       18.5       Trp       W       TGG       100			000	V				100	0.5
Ile       I       ATT       50       ACG       18.9         ATC       48.5       ACG       6.9         ATA       1.5       Val       V       GTT       51.2         GTC       33.8       GTA       12.3         AAG       1       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100         TTA       18.5       Trp       W       TGG       100	His	н	CAT	78.8		Thr	T	ACT	39.2
Ile       I       ATT       50       ACG       6.9         ATC       48.5       48.5       6.9         ATA       1.5       Val       V       GTT       51.2         GTC       33.8       GTA       12.3         AAG       1       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100         TTA       18.5       TTP       W       TGG       100									. 35
Ile       I       ATT       50       ACG       6.9         ATC       48.5       ST       Val       V       GTT       51.2         GTC       33.8       GTA       12.3       GTA       12.3         Lys       K       AAA       99       GTA       12.3         AAG       I       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100         TTA       18.5       TTP       W       TGG       100								:	18.9
ATC 48.5 ATA 1.5 Val V GTT 51.2 GTC 33.8 Comparison of the comparison of the comp	Ile	1	ATT	50					
Lys     K     AAA     99     GTG     33.8       Lys     K     AAA     99     GTA     12.3       Leu     L     TTG     64.5     Trp     W     TGG     100       TTA     18.5	110	•							
Lys K AAA 99 GTA 12.3 GTG 2.7  Leu L TTG 64.5 Trp W TGG 100 TTA 18.5						Val	ν	GTT	51.2
Lys       K       AAA       99       GTA       12.3         AAG       1       GTG       2.7         Leu       L       TTG       64.5       Trp       W       TGG       100         TTA       18.5			7171				•		
AAG 1 GTG 2.7  Leu L TTG 64.5 Trp W TGG 100 TTA 18.5	Lvc	v	A A A	99					
Leu L TTG 64.5 Trp W TGG 100 TTA 18.5	Lys	K							
TTA 18.5		-	AAG					010	2.1
	Leu	L	TTG	64.5		Trp	W	TGG	100
CTC 8 Tyr Y TAT 62			TTA	18.5					
			CTC	8		Tyr	Y	TAT	62
CTT 6.5 TAC 38			CTT	6.5				TAC	38
CTG 2.5				2.5					
CTA 0 Stop TAA 75						Stop		TAA	<b>75</b>
TGA 25			<del></del> -			•			
Met M ATG 100 TAG 0	Met	М	ATG	100					
	1.101	474	2410					/	-
Asn N AAT 52	Asn	N	AAT	52				į.	
AAC 48				48					

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## Purification of Native Der p III

Using a method adapted from Heymann et al. (Heymann et al. J Allergy Clin Immunol (1989) 83: 1055-1067), 15 ml of a 50-80% saturated ammonium sulfate precipitate of D. pteronyssinus spent growth medium was applied to an upward flowing, 2 cm x 90 cm polyacrylamide P-100 column equilibrated in PBS (Pharmacia, LKB Biotechnology, Uppsala, Sweden). The protein content of the eluted 5ml fractions was determined by measuring optical density (A280nm) and analysis by SDS-PAGE. Those fractions containing predominantly bands in the 30kDa region were pooled, concentrated by polyethylene glycol 6000 (BDH Chemicals, Aust. PTY. Ltd. Kilsyth, Vic. Aust), dialyzed against PBS and passed over the column again. This was repeated twice until the analysis by SDS-PAGE determined that the only bands detectable were a doublet of approximately molecular weight 30kDa.

## Protein Sequence Analysis

The affinity purified native Der p III and Der f III proteins were subjected to HPEC using a 12% column system (Applied Biosystems, Foster City, CA). The affinity purified proteins or the HPEC fractions of the appropriate molecular weight were then subjected to protein sequence analysis using Applied Biosystems model 477A gas-phase sequenator with on-line phenylthiohydantoin derivative analysis (model 120). Alternatively, the affinity purified proteins were first separated by SDS-PAGE (BioRad), transferred to Problot (Applied Biosystems) and sequenced using a Beckman model LF3000. After initial sequence analysis of the proteins, o-phthalaldehyde was applied to block N-termini except those with prolines (position 13 for Der p III) to eliminate contaminating peptide sequences and extend the N-terminal sequence unambiguously. The CNBr peptides were produced by cleaving the affinity purified proteins with 2% (w/v) CNBr in 70% (w/v) formic acid overnight at room temperature. The digested peptides were subjected to HPEC for purification and then the peptide fragments were subjected to sequence analysis.

## Preferred Codon Usage

The codon usage bias for the mature proteins Der f I (Dilworth et al. Clin Exp Allergy (1991) 21:25-32), Der f II (Trudinger et al. Clin Exp Allergy (1991) 21:33-37), Der p I (Chua et al. J Exp Med (1988) 167:175-182) and Der p II (Chua et al. Int Arch Allergy Appl Immunol (1990) 91:118-123) was determined. The average percent usage of each triplet codon for each amino acid was determined for each of these four Dermatophagoides proteins and the results tabulated in Table 1.

mite). Such peptides can also be used to study the mechanism of immunotherapy of *D.* pteronyssinus allergy and to design modified derivatives or analogs useful in immunotherapy.

Work by others has shown that high doses of allergen extracts generally produce the best results during immunotherapy (i.e., best symptom relief). However, many subjects are unable to tolerate large doses of such extracts due to systemic reactions elicited by the allergens and other components within these preparations. A peptide having an activity of Der p III has the advantage of being free of all other mite protein. Thus, such a peptide can be administered for therapeutic purposes with fewer anticipated side effects.

It is now also possible to design an agent or a drug capable of blocking or inhibiting the ability of a house dust mite allergen to induce an allergic reaction in sensitive subjects. Such agents could be designed, for example, in such a manner that they would bind to relevant anti-Der p III IgE molecules, thus preventing IgE-allergen binding, and subsequent mast cell/basophil degranulation. Alternatively, such agents could bind to cellular components of the immune system, resulting in suppression or desensitization of the allergic responses to house dust mite allergens. A non-restrictive example of this is the use of peptides including B or T cell epitopes of Der p III or modifications thereof, based on the cDNA protein structure of Der p III to suppress the allergic response to a house dust mite allergen. This could be carried out by defining the structures of fragments encoding B and T cell epitopes which affect B and T cell function in in vitro studies with blood components from subjects sensitive to house dust mite.

The invention is further illustrated by the following examples which should not be construed as further limiting the subject invention. The contents of all references and published patent applications cited throughout this application are hereby incorporated by reference.

The following methodology was used throughout the Examples.

#### **EXAMPLES**

#### 30 Materials and Methods

#### D. pteronyssinus Cultures

Whole mites were purchased from the Commonwealth Serum Laboratories, Parkville, Australia and spent medium was a gift from the same source.

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It is especially advantageous to formulate parenteral compositions in dosage unit form for ease of administration and uniformity of dosage. Dosage unit form as used herein refers to physically discrete units suited as unitary dosages for the mammalian subjects to be treated; each unit containing a predetermined quantity of active compound calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier. The specification for the dosage unit forms of the invention are dictated by and directly dependent on (a) the unique characteristics of the active compound and the particular therapeutic effect to be achieved, and (b) the limitations inherent in the art of compounding such an active compound for the treatment of sensitivity in subjects.

The invention also provides a composition comprising at least two peptides having an activity of Der p III (e.g., a physical mixture of at least two peptides), each having T cell stimulating activity. Alternatively, a peptide having at least two regions, each having T cell stimulating activity (i.e., each region comprising at least one T cell epitope) can be administered to an allergic subject. A composition of two peptides having a Der p III activity or a composition of two peptides having at least two regions, each having T cell stimulating activity can be administered to a subject in the form of a composition with a pharmaceutically acceptable carrier as hereinbefore described. An amount of one or more of such compositions can be administered simultaneously or sequentially to a subject sensitive to a house dust mite allergen to treat such sensitivity.

The cDNA (or the mRNA which served as a template during reverse transcription) encoding a peptide having an activity of Der p III can be used to identify similar nucleic gacids in any variety or type of animal and, thus, to molecularly clone genes which have sufficient sequence homology to hybridize to the cDNA encoding a peptide having an activity of Der p III. Thus, the present invention includes not only peptides having an activity of Der p III, but also other proteins which may be allergens encoded by DNA which hybridizes to DNA of the present invention.

Isolated peptides that are immunologically related to Der p III, such as by antibody cross-reactivity or T cell cross-reactivity, other than those already identified, are within the scope of the invention. Such peptides bind antibodies specific for the protein and peptides of the invention, or stimulate T cells specific for the protein and peptides of this invention.

A peptide having an activity of Der p III (i.e., Der p III produced recombinantly or by chemical synthesis) is free of all other house dust mite proteins and, thus, is useful in the standardization of allergen extracts which are key reagents for the diagnosis and treatment of house dust mite hypersensitivity. In addition, such a peptide is of a consistent, well-defined composition and biological activity for use in preparations that will be administered for therapeutic purposes (e.g., to modify the allergic response of a subject sensitive to house dust WO 95/15976 PCT/US94/14073

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coating such as lecithin, by the maintenance of the required particle size in the case of dispersion and by the use of surfactants. Prevention of the action of microorganisms can be achieved by various antibacterial and antifungal agents, for example, parabens, - chlorobutanol, phenol, ascorbic acid, thimerosal, and the like. In many cases, it will be preferable to include isotonic agents, for example, sugars, sodium chloride, or polyalcohols such as mannitol and sorbitol, in the composition. Prolonged absorption of the injectable compositions can be brought about by including in the composition an agent which delays absorption, for example, aluminum monostearate or gelatin.

Sterile injectable solutions can be prepared by incorporating active compound (i.e., at least one peptide having an activity of Der p III) in the required amount in an appropriate solvent with one or a combination of ingredients enumerated above, as required, followed by filtered sterilization. Generally, dispersions are prepared by incorporating the active compound into a sterile vehicle which contains a basic dispersion medium and the required other ingredients from those enumerated above. In the case of sterile powders for the preparation of sterile injectable solutions, the preferred methods of preparation are vacuum drying and freeze-drying which yields a powder of the active ingredient (i.e., at least one peptide having an activity of Der p III) plus any additional desired ingredient from a previously sterile-filtered solution thereof.

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When the peptide having an activity of Der p III is suitably protected, as described above, the peptide may be orally administered, for example, with an inert diluent or an assimilable edible carrier. The peptide and other ingredients may also be enclosed in a hard or soft shell gelatin capsule, compressed into tablets, or incorporated directly into the subject's diet. For oral therapeutic administration, the active compound may be incorporated with excipients and used in the form of ingestible tablets, buccal tablets, troches, capsules, elixirs, suspensions, syrups, wafers, and the like. The percentage of the active ingredient in the compositions and preparations may, of course, be varied and may conveniently be between about 5 to about 80% of the weight of the unit. The amount of active compound in such therapeutically useful compositions is such that a suitable dosage will be obtained.

As used herein "pharmaceutically acceptable carrier" includes any and all solvents, dispersion media, coatings, antibacterial and antifungal agents, isotonic and absorption delaying agents, and the like. The use of such media and agents for pharmaceutically active substances is well known in the art. Except insofar as any conventional media or agent is incompatible with the active compound, use thereof in the compositions suitable for pharmaceutical administration is contemplated. Supplementary active compounds can also be incorporated into the compositions.

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provide the optimum therapeutic response. For example, several divided doses may be administered daily or the dose may be proportionally reduced as indicated by the exigencies of the therapeutic situation.

The active compound (i.e., at least one peptide having an activity of Der p III) may be administered in a convenient manner such as by injection (subcutaneous, intravenous, etc.), oral administration, inhalation, transdermal application, or rectal administration. Depending enon the route of administration, the active compound may be coated in a material to protect the compound from the action of enzymes, acids and other natural conditions which may inactivate the compound.

To administer a peptide having an activity of Der p III by other than parenteral 10 administration, it may be necessary to coat the peptide with, or co-administer the peptide with, a material to prevent its inactivation. For example, a peptide having an activity of Der p III may be administered to a subject in an appropriate carrier, diluent or adjuvant, coadministered with enzyme inhibitors or in an appropriate carrier such as liposomes. Pharmaceutically acceptable diluents include saline and aqueous buffer solutions. Adjuvant is used in its broadest sense and includes any immune stimulating compound such as interferon. Adjuvants contemplated herein include resorcinols, non-ionic surfactants such as polyoxyethylene oleyl ether and n-hexadecyl polyethylene ether. Enzyme inhibitors include pancreatic trypsin inhibitor, diisopropylfluorophosphate (DEP) and trasylol. Liposomes include water-in-oil-in-water CGF emulsions as well as conventional liposomes (Strejan et al., (1984) J. Neuroimmunol. 7:27). For purposes of inducing T cell non-responsiveness, the composition is preferably administered in non-immunogenic form, e.g., one that does not contain adjuvant.

The active compound may also be administered parenterally or intraperitoneally. Dispersions can also be prepared in glycerol, liquid polyethylene glycols, and mixtures thereof and in oils. Under ordinary conditions of storage and use, these preparations may contain a preservative to prevent the growth of microorganisms.

Pharmaceutical compositions suitable for injectable use include sterile aqueous solutions (where water soluble) or dispersions and sterile powders for the extemporaneous preparation of sterile injectable solutions or dispersion. In all cases, the composition must be sterile and must be fluid to the extent that easy syringability exists. It must be stable under the conditions of manufacture and storage and must be preserved against the contaminating action of microorganisms such as bacteria and fungi. The carrier can be a solvent or dispersion medium containing, for example, water, ethanol, polyol (for example, glycerol, propylene glycol, and liquid polyetheylene glycol, and the like), suitable mixtures thereof, and vegetable oils. The proper fluidity can be maintained, for example, by the use of a

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radioimmunoassays (RIA), immuno-radiometric assays (IRMA), luminescence immunoassays (LIA), histamine release assays and IgE immunoblots.

In another assay, the presence in a subject of IgE specific for Der p III and the ability . of T cells of the subject to respond to T cell epitopes of Der p III can be determined by administering to the subject an Immediate Type Hypersensitivity test and/or a Delayed Type Hypersensitivity test (see e.g., Immunology (1985) Roitt, I.M., Brostoff, J., Male, D.K. (eds.), C.V. Mosby Co., Gower Medical Publishing, London, NY, pp. 19.2-19.18; pp.22.1-22.10) utilizing a peptide having an activity of Der p III, or a modified form of a peptide having an activity of Der p III, each of which binds IgE specific for the allergen. The same subjects are administered a Delayed Type Hypersensitivity test prior to, simultaneously with, or subsequent to administration of the Immediate Type Hypersensitivity test. Of course, if the Immediate Type Hypersensitivity test is administered prior to the Delayed Type Hypersensitivity test, the Delayed Type Hypersensitivity test would be given to those subjects exhibiting a specific Immediate Type Hypersensitivity reaction. The Delayed Type Hypersensitivity test utilizes a peptide having an activity of Der p III which has human T cell stimulating activity and which does not bind IgE specific for the allergen in a substantial percentage of the population of subjects sensitive to the allergen (e.g., at least about 75%). Those subjects found to have both a specific Immediate type Hypersensitivity reaction and a specific Delayed Type Hypersensitivity reaction are administered an amount of a composition suitable for pharmaceutical administration. The composition comprises the peptide having an activity of Der p III as used in the Delayed Type Hypersensitivity test and a pharmaceutically acceptable carrier or diluent.

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A peptide having an activity of *Der p* III can be used in methods of treating and preventing allergic reactions to a house dust mite allergen or a cross-reactive protein allergen. Thus, the present invention provides compositions suitable for pharmaceutical administration comprising an amount of at least one peptide having an activity of *Der p* III and a pharmaceutically acceptable carrier. Administration of the compositions of the present invention to a subject to be desensitized can be carried out using known procedures, at dosages and for periods of time effective to reduce sensitivity (i.e., reduce the allergic response) of the subject to house dust mite. The term subject is intended to include living organisms in which an immune response can be elicited, e.g., mammals. Examples of subjects include humans, dogs, cats, mice, rats, and transgenic species thereof. An amount of at least one peptide having an activity of *Der p* III necessary to achieve a therapeutic effect may vary according to factors such as the degree of sensitivity of the subject to house dust mite, the age, sex, and weight of the subject, and the ability of a peptide having an activity of *Der p* III to elicit an antigenic response in the subject. Dosage regima may be adjusted to

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techniques described herein. Such antibodies can be used to block or interfere with the T cell interaction with peptides presented by MHC.

Exposure of allergic subjects to peptides having an activity of Der p III and which have T cell stimulating activity may cause the appropriate T cell subpopulations to become non-responsive to the respective protein allergen (e.g., fail to stimulate an immune response upon such exposure). In addition, such administration may modify the lymphokine secretion profile as compared with exposure to the naturally-occurring protein allergen or portion thereof (e.g., result in a decrease of IL-4 and/or an increase in IL-2). Furthermore, exposure to peptides having an activity of Der p III which have T cell stimulating activity may 10 influence T cell subpopulations which normally participate in the response to the allergen such that these T cells are drawn away from the site(s) of normal exposure to the allergen (e.g., nasal mucosa, skin, and lung) towards the site(s) of therapeutic administration of the protein or fragment derived therefrom. This redistribution of T cell subpopulations may ameliorate or reduce the ability of a subject's immune system to stimulate the usual immune response at the site of normal exposure to the allergen, resulting in a diminution in allergic symptoms.

A peptide having an activity of Der p III when administered to a subject sensitive to house dust mites is capable of modifying the B cell response, T cell response, or both the B cell and the T cell response of the subject to the allergen. As used herein, modification of the allergic response of a subject to a house dust mite allergen can be defined as nonresponsiveness or diminution in symptoms to the allergen, as determined by standard clinical procedures (See e.g., Varney et al., (1990) British Medical Journal 302:265-269), including diminution in house dust mite induced asthmatic symptoms. As referred to herein, a diminution in symptoms includes any reduction in the allergic response of a subject to the 25 allergen following a treatment regimen with a peptide of the invention. This diminution in symptoms may be determined subjectively (e.g., the subject feels more comfortable upon exposure to the allergen), or clinically, such as with a standard skin test.

Peptides or antibodies of the present invention can also be used for detecting and diagnosing sensitivity to Der p III. For example, this could be done by combining blood or blood products obtained from a subject to be assessed for sensitivity with a peptide having an activity of Der p III, under conditions appropriate for binding of components in the blood (e.g., antibodies, T cells, B cells) with the peptide(s) and determining the extent to which such binding occurs. Other diagnostic methods for allergic diseases which the peptides or antibodies of the present invention can be used include radio-allergosorbent test (RAST), paper radioimmunosorbent test (PRIST), enzyme linked immunosorbent assay (ELISA),

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<u>Cancer Therapy</u> (1985) Alan R. Liss, Inc. pp. 77-96). Hybridoma cells can be screened immunochemically for production of antibodies specifically reactive with a peptide having an activity of *Der p* III and the monoclonal antibodies isolated.

The term antibody as used herein is intended to include fragments thereof which are also specifically reactive with a peptide having an activity of  $Der\ p$  III. Antibodies can be fragmented using conventional techniques and the fragments screened for utility in the same manner as described above for whole antibodies. For example,  $F(ab')_2$  fragments can be generated by treating antibody with pepsin. The resulting  $F(ab')_2$  fragment can be treated to reduce disulfide bridges to produce Fab' fragments. The antibody of the present invention is further intended to include bispecific and chimeric molecules having an anti- $Der\ p$  III portion.

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Another aspect of this invention provides T cell clones and soluble T cell receptors specifically reactive with a peptide having an activity of Der p III. Monoclonal T cell populations (i.e., T cells genetically identical to one another and expressing identical T cell receptors) can be derived from a subject sensitive to Der p III, followed by repetitive in vitro stimulation with a Der p III protein or peptide having an activity of Der p III in the presence of MHC-matched antigen-presenting cells. Single Der p III MHC responsive cells can then be cloned by limiting dilution and permanent lines expanded and maintained by periodic in vitro restimulation. Alternatively, Der p III specific T-T hybridomas can be produced by a technique similar to B cell hybridoma production. For example, a mammal, such as a mouse, is immunized with a peptide having an activity of Der p III, T cells from the mammal can be purified and fused with an autonomously growing T cell tumor line. From the resulting hybridomas, cells responding to a peptide having an activity of Der p III are selected and cloned. Procedures for propagating monoclonal T cell populations are described in Cellular and Molecular Immunology (Abul K. Abbas et al. ed.), W.B. Saunders Company, Philadelphia, PA (1991) page 139. Soluble T cell receptors specifically reactive with a peptide having an activity of Der p III can be obtained by immunoprecipitation using an antibody against the T cell receptor as described in Immunology: A Synthesis (Second Edition), Edward S. Golub et al., ed., Sinauer Associates, Inc., Sunderland, MA (1991) pages 366-369.

T cell clones specifically reactive with a peptide having an activity of Der p III can be used to isolate and molecularly clone the gene encoding the relevant T cell receptor. In addition, a soluble T cell receptor specifically reactive with a peptide having an activity of Der p III can be used to interfere with or inhibit antigen-dependent activation of the relevant T cell subpopulation, for example, by administration to a subject sensitive to Der p III. Antibodies specifically reactive with such a T cell receptor can be produced according to the

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acid pairs, such as KK or RR, can be introduced between regions within a protein or fragment during recombinant construction thereof. The resulting peptide can be rendered sensitive to cleavage by cathepsin and/or other trypsin-like enzymes which would generate portions of the protein containing one or more T cell epitopes. In addition, such charged amino acid residues can result in an increase in the solubility of the peptide.

Site-directed mutagenesis of a nucleic acid encoding a peptide having an activity of Der p III can be used to modify the structure of the peptide by methods known in the art. Such methods may, among others, include polymerase chain reaction (PCR) with oligonucleotide primers bearing one or more mutations (Ho et al., (1989) Gene 77:51-59) or total synthesis of mutated genes (Hostomsky, Z. et al., (1989) Biochem. Biophys. Res. Comm 161:1056-1063). To enhance recombinant protein expression, the aforementioned methods can be applied to change the codons present in the cDNA sequence of the invention to those preferentially utilized by the host cell in which the recombinant protein is being expressed (Wada et al., supra).

Another aspect of the invention pertains to an antibody specifically reactive with a peptide having an activity of Der p III. The antibodies of this invention can be used to standardize allergen extracts or to isolate the naturally-occurring or native form of Der p III. For example, by using peptides having an activity of Der p III based on the cDNA sequence of Der p III, anti-protein/anti-peptide antisera or monoclonal antibodies can be made using standard methods. A mammal such as a mouse, a hamster or a rabbit can be immunized with an immunogenic form of the peptide (e.g., Der p III protein or an antigenic fragment which is capable of eliciting an antibody response). Techniques for conferring immunogenicity on a protein or peptide include conjugation to carriers or other techniques are well known in the art. A peptide having an activity of Der p III can be administered in the presence of 25 adjuvant. The progress of immunization can be monitored by detection of antibody titers in plasma or serum. Standard ELISA or other immunoassays can be used with the immunogen as antigen to assess the levels of antibodies.

Following immunization, anti-Der p III antisera can be obtained and, if desired, polyclonal anti-Der p III antibodies isolated from the serum. To produce monoclonal antibodies, antibody-producing cells (lymphocytes) can be harvested from an immunized animal and fused by standard somatic cell fusion procedures with immortalizing cells such as myeloma cells to yield hybridoma cells. Such techniques are well known in the art, for example the hybridoma technique originally developed by Kohler and Milstein, (Nature (1975) 256:495-497) as well as other techniques such as the human B cell hybridoma technique (Kozbar et al., Immunology Today (1983) 4:72) and the EBV-hybridoma technique to produce human monoclonal antibodies (Cole et al., Monoclonal Antibodies and

acid residues which are not essential for interaction with the MHC protein complex but which still bind the MHC protein complex can be modified by being replaced by another amino acid whose incorporation may enhance, not affect, or diminish but not eliminate T cell reactivity. Preferred amino acid substitutions for non-essential amino acids include, but are not limited to substitutions with alanine, glutamic acid, or a methyl amino acid.

Another example of modification of a peptide having an activity of *Der p* III is substitution of cysteine residues preferably with alanine, serine, threonine, leucine or glutamic acid residues to minimize dimerization via disulfide linkages. In addition, amino acid side chains of peptides of the invention can be chemically modified. Another modification is cyclization of the peptide.

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In order to enhance stability and/or reactivity, a peptide having an activity of *Der p* III can be modified to incorporate one or more polymorphisms in the amino acid sequence of the protein allergen resulting from any natural allelic variation. Additionally, D-amino acids, non-natural amino acids, or non-amino acid analogs can be substituted or added to produce a modified protein within the scope of this invention. Furthermore, a peptide having an activity of *Der p* III can be modified using polyethylene glycol (PEG) according to the method of A. Sehon and co-workers (Wie et al., supra) to produce a protein conjugated with PEG. In addition, PEG can be added during chemical synthesis of the protein. Other modifications of a peptide having an activity of *Der p* III include reduction/alkylation (Tarr, Methods of Protein Microcharacterization, J. E. Silver ed., Humana Press, Clifton NJ 155-194 (1986)); acylation (Tarr, supra); chemical coupling to an appropriate carrier (Mishell and Shiigi, eds., Selected Methods in Cellular Immunology, WH Freeman, San Francisco, CA (1980), U.S. Patent 4,939,239; or mild formalin treatment (Marsh (1971), Int. Arch. of Allergy and Appl. Immunol. 41:199-215).

To facilitate purification and potentially increase solubility of a peptide having an activity of *Der p* III, it is possible to add an amino acid fusion moiety to the peptide backbone. For example, hexa-histidine can be added to the protein for purification by immobilized metal ion affinity chromatography (Hochuli, E. et al., (1988) Bio/Technology 6:1321-1325). In addition, to facilitate isolation of peptides free of irrelevant sequences, specific endoprotease cleavage sites can be introduced between the sequences of the fusion moiety and the peptide. In order to successfully desensitize a subject to *Der p* III protein or related allergen, it may be necessary to increase the solubility of the protein by adding functional groups to the protein or by omitting hydrophobic regions of the protein.

To potentially aid proper antigen processing of T cell epitopes within Der p III, canonical protease sensitive sites can be engineered between regions, each comprising at least one T cell epitope via recombinant or synthetic methods. For example, charged amino

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volume of whole heparinized blood from an allergic subject. After mixing and incubation, the cells are pelleted and the supernatants are processed and analyzed using a radioimmunoassay to determine the amount of histamine released.

Peptides having an activity of Der p III which are to be used as therapeutic agents are preferably tested in mammalian models of house dust mite atopy, such as the mouse model disclosed in Tamura et al., (1986) Microbiol. Immunol. 30:883-896, or in U.S. Patent 4,939,239, or in the primate model disclosed in Chiba et al., (1990) Int. Arch. Allergy Immunol. 93:83-88. Initial screening for IgE binding to a peptide having an activity of Der p III may be performed by scratch tests or intradermal skin tests on laboratory animals or human volunteers, or in in vitro systems such as RAST, RAST inhibition, ELISA assay, RIA (radioimmunoassay), or a histamine release assay, as described above.

It is possible to modify the structure of a peptide having an activity of Der p III for such purposes as increasing solubility, enhancing therapeutic or prophylactic efficacy, or stability (e.g., shelf life ex vivo and resistance to proteolytic degradation in vivo). Such modified peptides are considered functional equivalents of peptides having an activity of Der p III as defined herein. A modified peptide can be produced in which the amino acid sequence has been altered, such as by amino acid substitution, deletion, or addition, to modify immunogenicity and/or reduce allergenicity, or to which a component has been added for the same purpose.

For example, a peptide having an activity of Der p III can be modified so that it maintains the ability to induce T cell non-responsiveness and bind MHC proteins without the ability to induce a strong proliferative response or, possibly, any proliferative response when administered in immunogenic form. In this instance, critical binding residues for T cell receptor function can be determined using known techniques (e.g., substitution of each residue and determination of the presence or absence of T cell reactivity). Those residues shown to be essential for interaction with the T cell receptor can be modified by replacing the essential amino acid with another, preferably similar amino acid residue (a conservative substitution) whose presence is shown to enhance, diminish but not eliminate, or not affect T cell reactivity. In addition, those amino acid residues which are not essential for T cell receptor interaction can be modified by being replaced by another amino acid whose incorporation may enhance, diminish but not eliminate, or not affect T cell reactivity, but whose incorporation does not eliminate binding to relevant MHC.

Additionally, a peptide having an activity of Der p III can be modified by replacing an amino acid shown to be essential for interaction with the MHC protein complex with another, preferably similar amino acid residue (conservative substitution) whose presence is shown to enhance, diminish but not eliminate or affect T cell activity. In addition, amino

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248:1349-1356, for assay systems that can be used as the basis for an assay in accordance with the present invention.

In yet another embodiment, peptides having a Der p III activity are identified by IgE binding activity. For therapeutic purposes, peptides of the invention preferably do not bind 5 IgE specific for a house dust mite allergen or bind such IgE to a substantially lesser extent than the purified, native house dust mite allergen binds such IgE. Reduced IgE binding activity refers to IgE binding activity that is less than that of purified, native Der p III protein. If a peptide having a Der p III activity is to be used as a diagnostic reagent, it is not necessary that the peptide have reduced IgE binding activity compared to the native Der p III allergen. 10 IgE binding activity of peptides can be determined, for example, by an enzyme-linked immunosorbent assay (ELISA) using, for example, sera obtained from a subject (i.e., an allergic subject) that has been previously exposed to the native Der p III allergen. Briefly, the peptide suspected of having a Der p III activity is coated onto wells of a microtiter plate. After washing and blocking the wells, antibody solution consisting of the plasma of an allergic subject who has been exposed to a peptide suspected of having a Der p III activity is incubated in the wells. The plasma is generally depleted of IgG before incubation. A labeled secondary antibody is added to the wells and incubated. The amount of IgE binding is then quantified and compared to the amount of IgE bound by a purified, native Der p III protein. Alternatively, the IgE binding activity of a peptide can be determined by Western blot analysis. For example, a peptide suspected of having a Der p III activity is run on a polyacrylamide gel using SDS-PAGE. The peptide is then transferred to nitrocellulose and subsequently incubated with sera from an allergic subject. After incubation with a labeled secondary antibody, the amount of IgE bound is then determined and quantified.

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Another assay which can be used to determine the IgE binding activity of a peptide is a competition ELISA assay. Briefly, an IgE antibody pool is generated by combining plasma from house dust mite allergic subjects that have been shown by direct ELISA to have IgE reactive with native Der p III. This pool is used in ELISA competition assays to compare IgE binding of native Der p III and a peptide suspected of having a Der p III activity. IgE binding for the native Der p III and a peptide suspected of having a Der p III activity is determined and quantified.

If a peptide having an activity of Der p III binds IgE and is to be used as a therapeutic agent, it is preferable that such binding does not result in the release of mediators (e.g., histamines) from mast cells or basophils. To determine whether a peptide which binds IgE results in the release of mediators, a histamine release assay can be performed using standard reagents and protocols obtained from Amac, Inc. (Westbrook, ME). Briefly, a buffered solution of a peptide suspected of having a Der p III activity is combined with an equal

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immune cells to the site of antigen/T cell interaction, and activation of the B cell cascade, leading to the production of antibodies. One isotype of these antibodies, IgE, is fundamentally important to the development of allergic symptoms and its production is influenced early in the cascade of events at the level of the T helper cell, by the nature of the lymphokines secreted. A T cell epitope is the basic element, or smallest unit of recognition by a T cell receptor, where the epitope comprises amino acids essential to receptor recognition. Amino acid sequences which mimic those of the T cell epitopes and which modify the allergic response to protein allergens are within the scope of this invention.

Screening peptides for those which retain a *Der p* III activity as described herein can be accomplished using one or more of several different assays. For example, *in vitro*, *Der p* III T cell stimulatory activity is assayed by contacting a peptide known or suspected of having a *Der p* III activity with an antigen presenting cell which presents appropriate MHC molecules in a T cell culture. Presentation of a peptide having a *Der p* III activity in association with appropriate MHC molecules to T cells in conjunction with the necessary costimulation has the effect of transmitting a signal to the T cell that induces the production of increased levels of cytokines, particularly of interleukin-2 and interleukin-4. The culture supernatant can be obtained and assayed for interleukin-2 or other known cytokines. For example, any one of several conventional assays for interleukin-2 can be employed, such as the assay described in <u>Proc. Natl. Acad. Sci USA</u>, 86:1333 (1989), the pertinent portions of which are incorporated herein by reference. A kit for an assay for the production of interferon is also available from Genzyme Corporation (Cambridge, MA).

Alternatively, a common assay for T cell proliferation entails measuring tritiated thymidine incorporation. The proliferation of T cells can be measured *in vitro* by determining the amount of <sup>3</sup>H-labeled thymidine incorporated into the replicating DNA of cultured cells. Therefore, the rate of DNA synthesis and, in turn, the rate of cell division can be quantified.

In another embodiment, a peptide having a *Der p* III activity is screened for the ability to induce T cell non-responsiveness. The ability of a peptide known to stimulate T cells (as determined by one or more of the above described assays) to inhibit or completely block the activity of purified native *Der p* III or portion thereof and induce a state of non-responsiveness can be determined using subsequent attempts at stimulation of the T cells with antigen presenting cells that present native *Der p* III, or a peptide having a *Der p* III activity, following exposure to the peptide having a *Der p* III activity. If the T cells are unresponsive to the subsequent activation attempts, as determined by interleukin-2 synthesis and/or T cell proliferation, a state of non-responsiveness has been induced. See, e.g., Gimmi, et al. (1993) Proc. Natl. Acad. Sci USA, 90:6586-6590; and Schwartz (1990) Science,

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MICGGDVANGGVDSEQGD (SEQ ID NO:10), an internal peptide; and MTLDQTNAKPVPLPTS (SEQ ID NO:12), an internal peptide.

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Yet another embodiment of the invention provides a substantially pure preparation of a peptide having an activity of *Der p* III. Such a preparation is substantially free of proteins and peptides with which the peptide naturally occurs (i.e., other house dust mite peptides), either in a cell or when secreted by a cell.

The term isolated as used herein refers to a nucleic acid or peptide that is substantially free of cellular material or culture medium when produced by recombinant DNA techniques, or chemical precursors or other chemicals when chemically synthesized. Such peptides are also characterized as being free of all other house dust mite proteins. Accordingly, an isolated peptide having an activity of *Der p* III is produced recombinantly or synthetically and is substantially free of cellular material and culture medium or substantially free of chemical precursors or other chemicals and is free of all other house dust mite proteins. An isolated nucleic acid is also free of sequences which naturally flank the nucleic acid (i.e., sequences located at the 5' and 3' ends of the nucleic acid) in the organism from which the nucleic acid is derived.

Peptides having an activity of *Der p* III can be obtained, for example, by screening peptides recombinantly produced from the corresponding fragment of the nucleic acid of *Der p* III encoding such peptides. In addition, fragments can be chemically synthesized using techniques known in the art such as conventional Merrifield solid phase f-Moc or t-Boc chemistry. For example, the *Der p* III protein may be arbitrarily divided into fragments of desired length with no overlap of the fragments, or preferably divided into overlapping fragments of a desired length. The fragments can be produced (recombinantly or by chemical synthesis) and tested to identify those peptides having a *Der p* III activity, i.e., the ability to induce a T cell response such as T cell stimulation (T cell proliferation, cytokine secretion) or T cell non-responsiveness, and/or has reduced IgE binding activity.

In one embodiment, peptides having an activity of *Der p* III can be identified by the ability of the peptide to stimulate T cells or to induce T cell non-responsiveness. Peptides which stimulate T cells, as determined by, for example, T cell proliferation or cytokine secretion are defined herein as comprising at least one T cell epitope. T cell epitopes are believed to be involved in initiation and perpetuation of the immune response to the protein allergen which is responsible for the clinical symptoms of allergy. These T cell epitopes are thought to trigger early events at the level of the T helper cell by binding to an appropriate HLA molecule on the surface of an antigen presenting cell, thereby stimulating the T cell subpopulation with the relevant T cell receptor for the epitope. These events lead to T cell proliferation, lymphokine secretion, local inflammatory reactions, recruitment of additional

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including ion-exchange chromatography, gel filtration chromatography, ultrafiltration, electrophoresis, and immunoaffinity purification with antibodies specific for such peptides.

Another aspect of the invention pertains to isolated peptides having an activity of Der p III. A peptide having an activity of Der p III has at least one biological activity of the Der p III allergen. For example, a peptide having an activity of Der p III may have the ability to induce a response in Der p III specific T cells such as stimulation (T cell proliferation or cytokine secretion) or induce T cell non-responsiveness. In one embodiment, a peptide having an activity of Der p III stimulates T cells as evidenced by, for example, T cellproliferation or cytokine secretion. In another embodiment, peptides having a Der p III activity induce T cell non-responsiveness in which T cells are unresponsive to a subsequent challenge with a Der p III peptide or native Der p III protein following exposure to the peptide. In yet another embodiment, a peptide having a Der p III activity has reduced IgE binding activity compared to purified, native Der p III protein. A peptide having an activity of Der p III may differ in amino acid sequence from the Der p III sequence depicted in Figure 1A and 1B (SEQ ID NO:2) but such differences result in a modified protein which functions in the same or similar manner as a native Der p III protein or which has the same or similar characteristics of Der p III protein. Various modifications of the Der p III protein to produce these and other functionally equivalent peptides are described in detail herein. The term peptide, as used herein, refers to peptides, proteins, and polypeptides.

A peptide can be produced by modification of the amino acid sequence of the *Der p* III protein shown in Figure 1A and 1B (SEQ ID NO:2), such as by a substitution, addition, or deletion of one or more amino acid residues. Such modifications may be directed to amino acid residues not involved in a biological activity of the peptide, or the modifications may be directed to such amino acid residues in order to enhance, or eliminate a particular biological activity. Peptides of the invention are at least about 8 amino acid residues in length, preferably about 10-20 amino acid residues in length and more preferably about 10-16 amino acid residues in length. Peptides having an activity of *Der p* III of at least about 30 amino acid residues in length, at least about 40 amino acid residues in length, at least about 100 amino acid residues in length, at least about 100 amino acid residues in length, and at least about 200 amino acid residues or more in length are also included within the scope of this invention.

In another embodiment of the invention, peptides from a related Group III allergen of the species *Dermatophagoides farinae*, *Der f* III, are provided. Such peptides have been isolated from a purified, native *Der f* III protein and comprise the sequences:

IVGGVKAKAGDSPYQISLQSSSHFXGGSILD (SEQ ID NO:15), an N-terminal sequence;

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which renders the recombinant fusion protein easily purifiable by metal chelate chromatography.

Inducible non-fusion expression vectors include pTrc (Amann et al., (1988) Gene 69:301-315) and pET11d (Studier et al., Gene Expression Technology: Methods in Enzymology 185, Academic Press, San Diego, California (1990) 60-89). While target gene expression relies on host RNA polymerase transcription from the hybrid trp-lac fusion promoter in pTrc, expression of target genes inserted into pET11d relies on transcription from the T7 gn10-lac 0 fusion promoter mediated by coexpressed viral RNA polymerase (T7 gn1). This viral polymerase is supplied by host strains BL21(DE3) or HMS174(DE3) from a resident λ prophage harboring a T7 gn1 under the transcriptional control of the lacUV 5 promoter.

One strategy to maximize recombinant *Der p* III expression in *E. coli* is to express the protein in a host bacteria with an impaired capacity to proteolytically cleave the recombinant protein (Gottesman, S., Gene Expression Technology: Methods in Enzymology 185, Academic Press, San Diego, California (1990) 119-128). Another strategy would be to alter the nucleic acid encoding *Der p* III to be inserted into an expression vector so that the individual codons for each amino acid would be those preferentially utilized in highly expressed *E. coli* proteins (Wada et al., (1992) Nuc. Acids Res. 20:2111-2118). Such alteration of nucleic acids of the invention can be carried out by standard DNA synthesis techniques.

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The nucleic acids of the invention can also be chemically synthesized using standard techniques. Various methods of chemically synthesizing polydeoxynucleotides are known, including solid-phase synthesis which, like peptide synthesis, has been fully automated in commercially available DNA synthesizers (See, e.g., Itakura et al. U.S. Patent No. 4,598,049; Caruthers et al. U.S. Patent No. 4,458,066; and Itakura U.S. Patent Nos. 4,401,796 and 4,373,071, incorporated by reference herein).

The present invention further pertains to methods of producing peptides that have an activity of  $Der\ p$  III. For example, a host cell transfected with a nucleic acid vector directing expression of a nucleotide sequence encoding a peptide having an activity of  $Der\ p$  III can be cultured under appropriate conditions to allow expression of the peptide to occur. The peptide may be secreted and isolated from a mixture of cells and medium containing the peptide. Alternatively, the peptide may be retained cytoplasmically and the cells harvested, lysed and the protein isolated. A cell culture includes host cells, media and other byproducts. Suitable media for cell culture are well known in the art. Peptides of the invention can be isolated from cell culture medium, host cells, or both using techniques known in the art for purifying proteins

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E. coli, insect cells (baculovirus), yeast, or mammalian cells such as Chinese hamster ovary cell (CHO). Other suitable host cells are referred to in Goeddel (1990), supra, or known to those skilled in the art.

Expression in eucaryotic cells such as mammalian, yeast, or insect cells can lead to partial or complete glycosylation and/or formation of relevant inter- or intra-chain disulfide bonds of a recombinant peptide product. Examples of vectors for expression in yeast S. cerivisae include pYepSec1 (Baldari. et al., (1987) Embo J. 6:229-234), pMFa (Kurjan and Herskowitz, (1982) Cell 30:933-943), pJRY88 (Schultz et al., (1987) Gene 54:113-123), and pYES2 (Invitrogen Corporation, San Diego, CA). Baculovirus vectors available for the second of the pythological second of the pyt expression of proteins in cultured insect cells (SF 9 cells) include the pAc series (Smith et al., (1983) Mol. Cell Biol. 3:2156-2165) and the pVL series (Lucklow, V.A., and Summers, M.D., (1989) Virology 170:31-39). Generally, COS cells (Gluzman, Y., (1981) Cell 23:175-182) are used in conjunction with such vectors as pCDM 8 (Aruffo, A. and Seed, B., (1987) Proc. Natl. Acad. Sci. USA 84:8573-8577) for transient amplification/expression in mammalian cells, while CHO (dhfr-Chinese Hamster Qvary) cells are used with vectors 15 such as pMT2PC (Kaufman et al. (1987), EMBO J. 6:187-195) for stable amplification/expression in mammalian cells. Vector DNA can be introduced into mammalian cells via conventional techniques such as calcium phosphate or calcium chloride co-precipitation, DEAE-dextran-mediated transfection, or electroporation. Suitable methods for transforming host cells can be found in Sambrook et al. (Molecular Cloning: A 20 Laboratory Manual, 2nd Edition, Cold Spring Harbor Laboratory press (1989)), and other laboratory textbooks.

Expression in procaryotes is most often carried out in *E. coli* with either fusion or non-fusion inducible expression vectors. Fusion vectors usually add a number of NH<sub>2</sub> terminal amino acids to the expressed target gene. These NH<sub>2</sub> terminal amino acids often are referred to as a reporter group. Such reporter groups usually serve two purposes: 1) to increase the solubility of the target recombinant protein; and 2) to aid in the purification of the target recombinant protein by acting as a ligand in affinity purification. Often, in fusion expression vectors, a proteolytic cleavage site is introduced at the junction of the reporter group and the target recombinant protein to enable separation of the target recombinant protein from the reporter group subsequent to purification of the fusion protein. Such enzymes, and their cognate recognition sequences, include Factor Xa, thrombin and enterokinase. Typical fusion expression vectors include pGEX (Amrad Corp., Melbourne, Australia), pMAL (New England Biolabs, Beverly, MA) and pRIT5 (Pharmacia, Piscataway, NJ) which fuse glutathione S-transferase, maltose E binding protein, or protein A,

NJ) which fuse glutathione S-transferase, maltose E binding protein, or protein A, respectively, to the target recombinant protein. A preferred reporter group is poly(His),

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III. Generally, the nucleic acid encoding a peptide having an activity of Der p III will be selected from the bases encoding the mature protein, however, in some instances, it may be desirable to select all or part of a peptide from the leader sequence portion of the nucleic acid.
of the invention. Nucleic acids within the scope of the invention may also contain linker sequences, modified restriction endonuclease sites and other sequences useful for molecular cloning, expression or purification of recombinant peptides having an activity of Der p III.

A nucleic acid encoding a peptide having an activity of *Der p* III may be obtained from mRNA present in house dust mites of the species *Dermatophagoides pteronyssinus*. It should also be possible to obtain nucleic acids encoding *Der p* III from *Dermatophagoides pteronyssinus* genomic DNA. For example, the gene encoding *Der p* III can be cloned from either a cDNA or a genomic library in accordance with protocols herein described. A cDNA encoding *Der p* III can be obtained by isolating total mRNA from *Dermatophagoides pteronyssinus*. Double stranded cDNAs can then be prepared from the total mRNA. Subsequently, the cDNAs can be inserted into a suitable plasmid or bacteriophage vector using any one of a number of known techniques. Genes encoding *Der p* III can also be cloned using established polymerase chain reaction techniques in accordance with the nucleotide sequence information provided by the invention. The nucleic acids of the invention can be DNA or RNA. A preferred nucleic acid is a cDNA encoding *Der p* III having the sequence depicted in Figure 1A and 1B (SEQ ID NO:1).

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This invention also provides expression vectors containing a nucleic acid encoding a peptide having an activity of *Der p* III operably linked to at least one regulatory sequence. Operably linked is intended to mean that the nucleotide sequence is linked to a regulatory sequence in a manner which allows expression of the nucleotide sequence. Regulatory sequences are art-recognized and are selected to direct expression of the peptide having an activity of *Der p* III by a transfected host cell. Accordingly, the term regulatory sequence includes promoters, enhancers and other expression control elements. Such regulatory sequences are described in Goeddel, *Gene Expression Technology*: Methods in Enzymology 185, Academic Press, San Diego, CA (1990). It should be understood that the design of the expression vector may depend on such factors as the choice of the host cell to be transformed and/or the type of protein desired to be expressed. In one embodiment, the expression vector includes a DNA encoding a peptide having an activity of *Der p* III. Such expression vectors can be used to transfect cells to thereby produce proteins or peptides, including fusion proteins or peptides encoded by nucleic acids as described herein.

This invention further pertains to a host cell transfected to express a peptide having an activity of Der p III. The host cell may be any procaryotic or eucaryotic cell. For example, a peptide having an activity of Der p III may be expressed in bacterial cells such as

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stringency conditions at room temperature, about 22°C, to high stringency conditions at about 65°C.

Isolated nucleic acids encoding a peptide having an activity of Der p III, as described herein, and having a sequence that differs from the nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1) due to degeneracy in the genetic code are also within the scope of the invention. Such nucleic acids differ in sequence from the sequence of Figure 1A and 1B (SEQ ID NO:1) encode functionally equivalent peptides (i.e., peptides having an activity of Der p III), but due to degeneracy in the genetic code. For example, a number of amino acids are designated by more than one triplet. Codons that specify the same amino acid, or synonyms (for example, CAU and CAC are synonyms for histidine) may result in "silent" mutations that do not affect the amino acid sequence of the Der p III protein. However, it is expected that DNA sequence polymorphisms that do lead to changes in the amino acid sequence of Der p III will exist within the house dust mite population. One skilled in the art will appreciate that these variations in one or more nucleotides (up to about 3-4% of the nucleotides) of the nucleic acids encoding peptides having an activity of Der p III may exist among individual house dust mites due to natural allelic variation. Any and all such nucleotide variations and resulting amino acid polymorphisms are within the scope of this invention. Furthermore, there may be one or more isoforms or related, cross-reacting family members of Der p III. Such isoforms or family members are defined as proteins related in function and amino acid sequence to Der p III, but are encoded by genes at different loci.

Fragments of a nucleic acid encoding Der p III are also within the scope of the invention. As used herein, a fragment of a nucleic acid encoding Der p III refers to a nucleotide sequence having fewer nucleotides than the nucleotide sequence encoding the entire amino acid sequence of Der p III protein and which encodes a peptide having an activity of Der p III (i.e., a peptide having at least one biological activity of the Der p III allergen) as defined herein.

Preferred nucleic acid fragments encode peptides of at least 10 amino acid residues in length, preferably about 10-20 amino acid residues in length, and more preferably about 10-16 amino acid residues in length. Nucleic acid fragments that encode peptides having a Der p III activity of at least about 30 amino acid residues in length, at least about 40 amino acid residues in length, at least about 60 amino acid residues in length, at least about 80 amino acid residues in length, at least about 100 amino acid residues in length, and at least about 200 amino acid residues in length or more are also within the scope of this invention.

Nucleic acid fragments within the scope of the invention include those capable of hybridizing under high or low stringency with nucleic acids from other animal species for use in screening protocols to detect Der p III or allergens that are cross-reactive with Der p

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cytokine secretion) or induce T cell non-responsiveness. Alternatively, or additionally, a peptide having an activity of *Der p* III may have the ability to bind (to be recognized by) immunoglobulin E antibodies of house dust mite-allergic subjects. Peptides which bind IgE are useful in methods of detecting allergic sensitivity to *Der p* III in a subject. Peptides that do not bind IgE, or bind IgE to a lesser extent than a purified, native *Der p* III protein binds IgE are particularly useful as therapeutic agents.

In one embodiment, the nucleic acid is a cDNA encoding a peptide having an activity of *Der p* III. Preferably, the nucleic acid is a cDNA molecule comprising at least a portion of the nucleotide sequence encoding *Der p* III shown in Figure 1A and 1B (SEQ ID NO:1). A preferred portion of the cDNA molecule of Figure 1A and 1B (SEQ ID NO:1) includes the coding region of the molecule.

In another embodiment, the nucleic acid of the invention encodes a peptide having an activity of Der p III and comprising an amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2). Preferred nucleic acids encode a peptide having a Der p III activity and having at least about 50% homology, more preferably at least about 60% homology and most preferably at least about 70% homology with the sequence shown in Figure 1A and 1B (SEQ ID NO:2). Nucleic acids which encode peptides having a Der p III activity and having at least about 90%, more preferably at least about 95%, and most preferably at least about 98-99% homology with a sequence set forth in Figure 1A and 1B (SEQ ID NO:2) are also within the scope of the invention. Homology refers to sequence similarity between two peptides having an activity of Der p III or between two nucleic acid molecules. Homology can be determined by comparing a position in each sequence which may be aligned for purposes of comparison. When a position in the compared sequence is occupied by the same base or amino acid, then the molecules are homologous at that position. A degree of homology between sequences is a function of the number of matching or homologous positions shared by the sequences.

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Another aspect of the invention provides a nucleic acid that hybridizes under high or low stringency conditions to a nucleic acid encoding a peptide having all or a portion of an amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2). Appropriate stringency conditions which promote DNA hybridization, for example, 6.0 x sodium chloride/sodium citrate (SSC) at about 45°C, followed by a wash of 2.0 x SSC at 50° are known to those skilled in the art or can be found in <u>Current Protocols in Molecular Biology</u>, John Wiley & Sons, N.Y. (1989), 6.3.1-6.3.6. For example, the salt concentration in the wash step can be selected from a low stringency of about 2.0 x SSC at 50°C to a high stringency of about 0.2 x SSC at 50°C. In addition, the temperature in the wash step can be increased from low

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## **Detailed Description of the Invention**

This invention pertains to isolated nucleic acids encoding peptides having at least one biological activity of Der p III, a Group III allergen of the species Dermatophagoides pteronyssinus. Preferably, the nucleic acid is a cDNA comprising a nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1).

The cDNA shown in Figure 1A and 1B (SEQ ID NO:1) encodes a Der p III peptide which includes a predicted 29 amino acid residue pre-pro region encoded by nucleotides 63 through 149. This leader sequence is not found in the mature Der p III which is encoded by nucleotides 150 through 845. The deduced amino acid sequence based on this cDNA is also shown in Figure 1A and 1B (SEQ ID NO:2). The cDNA encodes a 232 residue mature peptide having a predicted molecular weight of 24,985 daltons including seven cysteine residues. A polyadenylation signal sequence 179 nucleotides after the last base is present in the cDNA (See Figure 1A and 1B). A culture of E. coli transfected with an expression vector containing the cDNA encoding Der p III was deposited under the Budapest Treaty with the American Type Culture Collection on October 15, 1993 and assigned accession number 69472.

Accordingly, one aspect of this invention pertains to isolated nucleic acids comprising a nucleotide sequence encoding Der p III, fragments thereof encoding peptides having at least one biological activity of Der p III and/or equivalents of such nucleic acids. The term nucleic acid as used herein is intended to include such fragments and equivalents. The term equivalent is intended to include nucleotide sequences encoding functionally equivalent Der p III peptides having an activity of Der p III. As defined herein, a peptide having an activity of Der p III has at least one biological activity of the Der p III allergen. Equivalent nucleotide sequences include sequences that differ by one or more nucleotide substitutions, additions or deletions, such as allelic variants, and include sequences that differ from the nucleotide sequence encoding Der p III shown in Figure 1A and 1B (SEQ ID NO:1) due to the degeneracy of the genetic code. Equivalents includes nucleotide sequences that hybridize under stringent conditions (i.e., equivalent to about 20-27°C below melting temperature (T<sub>m</sub>) and about 1M NaCl) to the nucleotide sequence of Der p III shown in Figure 1A and 1B (SEQ ID NO:1).

Peptides referred to herein as having an activity of Der p III or having a Der p III activity are defined herein as peptides that have an amino acid sequence corresponding to all or a portion of the amino acid sequence of Der p III shown in Figure 1A and 1B (SEQ ID NO:2). For example, a peptide having an activity of Der p III may have the ability to induce a response in Der p III restricted T cells such as stimulation (e.g., T cell proliferation or

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Peptides having a *Der p* III activity produced by recombinant expression of a nucleic acid of the invention, and peptides having a *Der p* III activity prepared by chemical synthesis are also featured by this invention. Preferred peptides have the ability to induce a T cell response which can include T cell stimulation (measured by, for example, T cell proliferation or cytokine secretion) or T cell non-responsiveness (i.e., contact with the peptide or a complex of the peptide with an MHC molecule of an antigen presenting cell induces the T cell to become unresponsive to stimulatory signals or incapable of proliferation). Other preferred peptides either apart from or in addition to the ability to induce a T cell response, have the ability to bind specific IgE of house dust mite-allergic subjects. Such peptides are useful in diagnosing sensitivity to house dust mites in a subject. Still other peptides, either apart from or in addition to the ability to induce a T cell response, have a significantly reduced or negligible ability to bind house dust mite-allergic IgE. Such peptides are particularly useful as therapeutic agents.

Other preferred peptides comprise an amino acid sequence shown in Figure 1A and 1B (SEQ ID NO:2). In one embodiment, peptides having a *Der p* III activity and comprising a portion of the amino acid sequence of Figure 1A and 1B (SEQ ID NO:2) are at least about 8-30 amino acids in length, preferably about 10-20 amino acids in length, and most preferably about 10-16 amino acids in length.

Another aspect of the invention features antibodies specifically reactive with peptides having a  $Der\ p$  III activity. Peptides having an activity of  $Der\ p$  III can be used in compositions suitable for pharmaceutical administration. For example, such compositions can be used in a manner similar to house dust mite extracts to treat or prevent allergic reactions to house dust mites in a subject. Nucleic acids of the invention and peptides having an activity of  $Der\ p$  III can also be used for diagnosing sensitivity in a subject to house dust mites.

#### **Brief Description of the Drawings**

Figure 1A and 1B show the complete nucleotide (SEQ ID NO:1) and deduced amino acid (SEQ ID NO:2) sequences of the *Der p* III clone.

Figure 2A and 2B show the amino acid sequences of *Der p* III (SEQ ID NO:2) and a trypsin protein from crayfish (SEQ ID NO:3).

Figure 3 shows the results of SDS-PAGE analysis of various concentrations of recombinant Der p III (lane 1, 4.3  $\mu$ g; lane 2, 8.7  $\mu$ g; lane 3, 13.8  $\mu$ g; lane 4, 17.4  $\mu$ g; markers are indicated by M).

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(1983) 17:263-73; Tovey E.R. et al., J. Allergy Clin. Immunol. (1987) 79:93-102) have shown that most sera contain IgE which recognize other allergens.

The significance of the IgE reactivity to *Der p* III remains uncertain. The reactivity of this group of allergens has been reported to be as low as 16% using a fluid phase assay (Heymann et al., J Allergy Clin Immunol (1989) 83: 1055-1067) and as high as 100% using RAST assay (Stewart et al., Immunology (1992) 75: 29-35). Several others have reported IgE reactivity between 60-83% (Tovey et al., J Allergy Clin Immunol (1987) 72: 93-102; Thomas et al., Exp Appl Acarol (1992) 16:153-164; Yasueda et al., Clin Exp Allergy (1993) 23:384-390). The discrepancies in the frequency of IgE reactivity to the group III allergens may be attributable to either the differences in the purity of the allergen preparation studied or the differences in sensitivity of the assay techniques used. In order to determine the importance of particular specificities in the allergic reactions, there is a need for quantities of pure allergen, which would enable quantitative IgE binding tests and studies of the frequency and lymphokine profile of T cell responses to the allergen.

Many patients with sensitivity to house dust mite allergens are treated currently by administration of small, gradually increasing doses of house dust mite extracts. Use of these extracts has multiple drawbacks, including potential anaphylaxis during treatment and the necessity of continuing therapy, often for a period of several years to build up sufficient tolerance and significant diminution of clinical symptoms. The ability to substitute compositions of house dust mite allergen, *Der p* III, would overcome several of these drawbacks. Thus, a source of pure allergen that could be provided in quantity for use as a diagnostic or therapeutic reagent and therapeutic methods that would overcome the drawbacks associated with house dust mite extracts are highly desirable.

#### **Summary of Invention**

This invention provides isolated nucleic acids encoding peptides having at least one biological activity of *Der p* III, an allergen of the species *Dermatophagoides pteronyssinus* (house dust mite). A preferred nucleic acid is a cDNA having a nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1). The invention also pertains to peptides encoded by all or a portion of such cDNA (SEQ ID NO:1) and having at least one biological activity of *Der p* III. Also contemplated are isolated nucleic acids which hybridize under high stringency conditions (e.g., equivalent to 20-27°C below T<sub>m</sub> and 1M NaCl ) to a nucleic acid having a nucleotide sequence shown in Figure 1A and 1B (SEQ ID NO:1) or which encodes a peptide comprising all or a portion of an amino acid sequence of Figure 1A and 1B (SEQ ID NO:2). Nucleic acids which encode peptides having an activity of *Der p* III and having at least 50% homology with a sequence shown in Figure 1A and 1B (SEQ ID NO:2) are also featured.

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# NUCLEIC ACIDS ENCODING A HOUSE DUST MITE ALLERGEN, DER P III, AND USES THEREFOR

#### **Background of the Invention**

Approximately 10% of the population become hypersensitized (allergic) upon exposure to antigens from a variety of environmental sources. Those antigens that induce immediate and/or delayed types of hypersensitivity are known as allergens (King, T.P., (1976) Adv. Immunol., 23:77-105). Allergens can include products of grasses, trees, weeds, animal dander, insects, food, drugs, and chemicals. Genetic predisposition of an individual is believed to play a role in the development of immediate allergic responses (Young, R. P. et al., (1990) Clin. Sci., 79:19) such as atopy and anaphylaxis, whose symptoms include hay fever, asthma, and hives.

The antibodies involved in atopic allergy belong primarily to the IgE class of immunoglobulins. IgE binds to basophils, mast cells and dendritic cells via a specific, highaffinity receptor FceRI (Kinet, J.P., (1990) Curr. Opin. Immunol., 2:499-505). Upon combination of an allergen acting as a ligand with its cognate receptor IgE, FceRI bound to the IgE may be cross-linked on the cell surface, resulting in physiological manifestations of the IgE - allergen interaction. These physiological effects include the release of, among other substances, histamine, serotonin, heparin, chemotactic factor(s) for eosinophilic leukocytes and/or leukotrienes C4, D4, and E4, which cause prolonged constriction of bronchial smooth muscle cells (Hood, L.E. et al., Immunology (2nd ed.), The Benjamin/Cumming Publishing Co., Inc. (1984)). Hence, the ultimate consequence of the interaction of an allergen with IgE is allergic symptoms triggered by the release of the aforementioned mediators. Such symptoms may be systemic or local in nature, depending on the route of entry of the antigen and the pattern of deposition of IgE on mast cells or basophils. Local manifestations generally occur on epithelial surfaces at the site of entry of the allergen. Systemic effects can induce anaphylaxis (anaphylactic shock) which results from IgE-basophil response to circulating (intravascular) antigen.

Studies with purified allergens have shown that about 80% of patients allergic to the mite *Dermatophagoides pteronyssinus* produce IgE reactive to *Der p* I and *Der p* II (Chapman M.D. et al., J. Immunol. (1980) 125:587-92; Lind P., J. Allergy Clin. Immunol. (1985) 76:753-61; Van der Zee J.S. et al., J. Allergy Clin. Immunol. (1988) 81:884-95). For about half the patients these specificities constitute 50% of the IgE antimite antibody. The allergen *Der p* III, recently identified as trypsin, (Stewart G.A. et al., Immunology (1992) 75:29-35) reacts with a similar or higher frequency (Stewart G.A. et al., supra; Ford S.A. et al., Clin. Exp. Allergy (1989) 20:27-31). However, in the only quantitative study reported to date, the investigators determined that the level of IgE binding to *Der p* III was considerably less than for *Der p* I. Electrophoretic techniques (Ford S.A. et al., supra; Bengtsson A. et al., Int. Arch. Allergy Appl. Immunol. (1986) 80:383-90; Lind P. et al., Scand. J. Immunol.

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6872 (AU).

Isolated nucleic acids encoding an allergen of Dermatophagoides pteronyssinus, Der p III, are disclosed. A cDNA encoding a peptide having a Der p III activity and a predicted molecular weight of about 24,985 daltons is also described. The nucleic acids can be used as probes to detect the presence of Der p III nucleic acid in a sample or for the recombinant production of peptides having an activity of Der p III. Peptides having an activity of Der p III can be used in compositions suitable for pharmaceutical administration or methods of diagnosing sensitivity to house dust mites.